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(71) Applicant: ADMINISTRATORS OF THE TULANE ED-UCATIONAL FUND [US/US]; 1430 Tulane Avenue, New Oleans, LA 70112 (US).

(72) Inventors: COY, David, H.; 4319 Perrier Street, New Orleans, LA 70115 (US). MOREAU, Jacques-Pierre; 159 Westboro Street, Upton, MA 01568 (US). TAYLOR, John, E.; 74 Fisk Hill Road, Upton, MA 01568 (US). KIM, Sun, Hyuk; 20 Whitney Street, Chestnut Hill, MA 02167 (US).

(74) Agent: KENNEDY, Bill; Fish & Richardson, One Financial Center, Suite 2500, Boston, MA 02111-2658 (US).

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(54) Title: THERAPEUTIC PEPTIDES

(57) Abstract

In general, the invention features a linear (i.e., non-cyclic) bombesin analog of biologically active mammalian gastrin-releasing peptide (GRP) and amphibian bombesin, having an active site and a binding site responsable for the binding of the peptide to a receptor on a target cell; cleavage of a peptide bond in the active site of naturally occurring bombesin or GRP is unnecessary for *in vivo* biological activity. The analog has one of the following modifications: (a) a deletion of a residue within the active site and a modification of a residue outside of the active site, or (b) a remplacement of one or two residues within the active site with a synthetic amino acid. The analog is capable of binding to the receptor and acting as a competitive inhibitor of the naturally occurring peptide by binding to the receptor and, by virtue of one of the modifications, failing to exhibit the *in vivo* biological activity of the naturally occurring peptide.

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THERAPEUTIC PEPTIDES

Background of the Invention

This invention relates to therapeutic peptides useful, e.g., for treatment of benign or malignant proliferation of tissue and for gastrointestinal disorders.

The amphibian peptide bombesin, pGlu-Gln-Arg-Leu-Gly-Asn-Gln-Trp-Ala-Val-Gly-His-Leu-Met-NH2 (Anastasi et al., Experientia 27:166-167 (1971)), is closely related to the mammalian gastrin-releasing peptides (GRP), e.g., the porcine GRP, H_2N- Ala-Pro-Val-Ser-Val-Gly-Gly-Gly-Thr-Val-Leu-Ala-Lys-Met-Tyr-Pro-Arg-Gly-Asn-His-Trp-Ala-Val-Gly-His-Leu-Met-(NH2) (McDonald et al., Biochem. 15 Biophys. Res. Commun. 90:227-233 (1979)) and human GRP, ${\tt H_2N-Val-Pro-Leu-Pro-Ala-Gly-Gly-Gly-Thr-Val-Leu-Thr-Lys-}$ Met-Tyr-Pro-Arg-Gly-Asn-His-Trp-Ala-Val-Gly-His-Leu-Met (NH_2) . Bombesin has been found to be a growth factor for a number of human cancer cell lines, including 20 small-cell lung carcinoma (SCLC), and has been detected in human breast and prostate cancer (Haveman et al., eds. <u>Recent Results in Cancer Research</u> - <u>Peptide</u> Hormones in Lung Cancer, Springer-Verlag, New York:1986). A number of these cancers are known to 25 secrete peptide hormones related to GRP or bombesin. Consequently, antagonists to bombesin have been proposed as agents for the treatment of these cancers.

Cuttitta et al. demonstrated that a specific monoclonal antibody to bombesin inhibited in vivo the growth of a human small-cell lung cancer cell line xenografted to nude mice (Cuttitta et al., Cancer Survey)

 $\underline{4}$:707-727 (1985)). In 3T3 murine fibroblasts which are responsive to the mitotic effect of bombesin, Zachary and Rozengurt observed that a substance P antagonist (Spantide) acted as a bombesin antagonist (Zachary et al., Proc. Natl. Acad. Sci. (USA), 82:7616-7620 (1985)). Heinz-Erian et al. replaced His at position 12 in bombesin with D-Phe and observed bombesin antagonist activity in dispersed acini from guinea pig pancreas (Heinz-Erian et al., Am. J. of Physiol. 252:G439-G442 10 (1987)). Rivier reported work directed toward restricting the conformational freedom of the bioactive C-terminal decapeptide of bombesin by incorporating intramolecular disulfide bridges; however, Rivier mentioned that, so far, bombesin analogs with this modification fail to exhibit any antagonist activity

15 (Rivier et al., "Competitive Antagonists of Peptide Hormones," in Abstracts of the International Symposium on Bombesin-Like Peptides in Health and Disease, Rome, Italy (October, 1987).

20 Abbreviations (uncommon):

pGlu = H, C--CH-COOH (pyroglutamatic acid);

 $Nle = H_2N-CH-COOH$ (norleucine) 25 (CH₂)₃-CH₃

Pal = 3-pyridyl-alanine

 β -leu = β - homoleucine

 γ -leu = gamma - homoleucine

30 D-Cpa = D-p-chlorophenylalanine

HyPro = hydroxyproline

Nal = naphthylalanine

Sar = sarcosine

- 3 -

Sta (statine) =

(3S, 4S)-4-amino-3-hydroxy-6-methylheptanoic acid, and has the chemical structure:

AHPPA =

(3S,4S)-4-amino-3-hydroxy-5-phenylpentanoic acid

10 ACHPA =

(3S, 4S)-4-amino-5-cyclohexyl-3-hydroxypentanoic acid; R = right (D) configuration; S = left (L) configuration; racemate = equal mix of R and S

1-methyl-His; 3-methyl-His = methyl (CH₃) group on nitrogen at positions 1 or 3 of Histidine:

$$CH_2-N^3H$$
 CH_2-N^1
 CH_2
 HCH
 $NH_2-CH-COOH$

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Summary of the Invention

A linear peptide (i.e., noncyclic) which is an analog of naturally occurring, biologically active amphibian bombesin or mammalian gastrin releasing peptide (GRP) having an active site and a binding site responsible for the binding of the peptide to a receptor on a target cell, cleavage of a peptide bond in the active site of naturally occurring bombesin or GRP being unnecessary for in vivo biological activity, the analog having one of the following modifications: (a) a deletion of a residue within the active site and a modification of a residue outside of the active site, or (b) a replacement of one or two residues within the

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active site with a synthetic amino acid.

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The analog is capable of acting as a competitive inhibitor of the naturally occurring peptide by binding to the receptor and, by virtue of one of the modifications, failing to exhibit the <u>in vivo</u> biological activity of the naturally occurring peptide.

The locations of the modifications that give rise to antagonists are determined by the location of the active site in the naturally occurring peptide. For example, the linear peptides for which introduction of a non-peptide bond between two residues, or the replacement of two natural amino acids with a synthetic β - or γ - amino acid, or the deletion ("des") of the C-terminal residue are useful in creating or enhancing antagonist activity are those in which activity is associated with the two C-terminal residues of the amino acid chain. Therefore, the active site of the naturally occurring peptide of which the peptides of the invention are analogs preferably includes at least one amino acid in the carboxy terminal half of the peptide, and the linear peptide of the invention includes that amino acid in its carboxy terminal half. Similarly, where the active site is located in the amino terminal portion of the naturally occurring peptide, the corresponding analogs of the invention will possess modifications in their amino terminal portions.

In preferred embodiments, the active site includes at least one amino acid residue located in the carboxyl terminal half of the naturally occurring biologically active peptide and that amino acid residue is located in the carboxyl terminal half of the linear peptide.

In other preferred embodiments, the active site includes at least one amino acid residue located in the

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amino terminal half of the naturally occurring biologically active peptide and that amino acid residue is located in the amino terminal half of the linear peptide.

Modifications can be introduced in a region involved in receptor binding, or in a non-binding region. Preferably, analogs of the invention are 25% homologous, most preferably, 50% homologous, with the naturally occurring peptides.

The analogs of the invention may have one of the modifications given in the generic formulae given below; either a non-peptide bond instead of a peptide bond between an amino acid of the active site and an adjacent amino acid; or a statine or AHPPA or ACHPA residue, or a β - or γ - amino acid in place of one or two natural amino acids; or a deletion of the C-terminal amino acid which may or may not be accompanied by the addition of a substituent on the actual C-terminal group or the presence of an N-terminal residue that is not the natural N-terminal amino acid of the peptides from which the analogs are derived. (Statine, AHPPA, and ACHPA have the chemical structures defined above. Where statine is used herein, AHPPA or ACHPA may also be used.)

By non-peptide bond is meant that the carbon atom participating in the bond between two residues is reduced from a carbonyl carbon to a methylene carbon, i.e., CH₂-NH; or, less preferably, that the residue bonded to the carbon atom has a sulfur atom or a methylene carbon in place of its amino group, i.e., CH₂-S or CO-CH₂. (A detailed discussion of the chemistry of non-peptide bonds is given in Coy et al. (1988) Tetrahedron 44,3:835-841, hereby incorporated by reference.)

One modification of the naturally occurring peptide to create an antagonist is of the amino terminal end of the molecule, which may be an electron-donating residue, e.g., a nitrogen-containing amino acid, such as those described for the amino terminal positions in the generic formulae below; for example, the N-terminal amino acid, which is A^0 or, if A^0 is deleted, is A^1 or, if A^0 and A^1 are deleted, is A^2 below, may be an aromatic D-isomer, or may be an alkylated amino acid. (Where "D" is not designated as the configuration of an amino acid, L is intended.) Another modification is of the C-terminal residue, which may be any of the "W" groups described below.

The invention includes a therapeutic peptide comprising between seven and nine amino acid residues, inclusive, the peptide being an analog of one of the following naturally occurring peptides terminating at the carboxy-terminus with a Met residue: (a) litorin; (b) neuromedin; (c) the ten amino acid carboxy-terminal region of mammalian gastrin releasing peptide; and (d) the ten amino acid carboxy-terminal region of amphibian bombesin, the therapeutic peptide being of the formula:

$$A^{0}-A^{1}-A^{2}-Trp-A^{4}-A^{5}-A^{6}-A^{7}-W$$

wherein

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A⁰ = Gly, Nle, α-aminobutyric acid, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β-Nal, or is deleted;

- 7 -

- A^1 = the D- or L-isomer of any of pGlu, Nle, or α -aminobutyric acid, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β -Nal, or is deleted;
- pGlu, Gly, Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, ß-Nal, His, l-methyl-His, or
- 10 A^4 = Ala, Val, Gln, Asn, Gly, Leu, Ile, Nle, α -aminobutyric acid, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β -Nal;
- $A^5 = Gln$, Asn, Gly, Ala, Leu, Ile, Nle, α -aminobutyric acid, Met, Val, p-X-Phe (where X = F, Cl, Br, OH, H or CH_3), Trp, Thr, or β -Nal;
 - A⁶ = Sar, Gly, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or B-Nal;
 - A' = 1-methyl-His, 3-methyl-His, or His; provided that if A^0 is present, A^1 cannot be pGlu; and if A^0 or A^1 is present, A^2 cannot be pGlu; and when A^0 is deleted and A^1 is pGlu, R_1 must be H and R_2 must be the portion of Glu that forms the imine ring in pGlu; and further provided that W can be:

3-methyl-His;

wherein R_3 is $CHR_{14}-(CH_2)_{n1}$ (where R_{14} is either H or OH; and n1 may be either of 1 or 0), or is deleted, and Z_1 is the identifying group of any one of

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the amino acids Gly, Ala, cyclohexyl-Ala, Val, Leu, Ile, Ser, Asp, Asn, Glu, Gln, p-X-Phe (where X = H, F, Cl, Br, NO₂, OH, or CH₃), Trp, Cys, Met, Pro, HyPro, or isopropyl, cyclohexylmethyl, β -nal, β -napthylmethyl, or phenylmethyl; and V is either

OR₄, or R₅

where R_4 is any of C_{1-20} alkyl, C_{3-20} alkenyl, C_{3-20} alkinyl, phenyl, napthyl, or C_{7-10} phenylalkyl, and each R_5 , and R_6 , independently, is any of H, C_{1-12} alkyl, C_{7-10} phenylalkyl, lower acyl, or R_{15}

where R₁₅ is any of H, C₁₋₁₂ alkyl, C₇₋₁₀ phenylalkyl, or lower acyl; provided that when one of R₅ or R₆ is NHR₁₅, the other is H; and provided that any asymmetric carbon atom can be R, S or a racemic mixture; and further provided that each R₁ and R₂, independently, is H, C₁₋₁₂ alkyl, C₇₋₁₀ phenylalkyl, COE₁ (where E₁ is C₁₋₂₀ alkyl, C₃₋₂₀ alkenyl, C₃₋₂₀ alkinyl, phenyl, naphthyl, or C₇₋₁₀ phenylalkyl), or lower acyl, and R₁ and R₂ are

bonded to the N-terminal amino acid of the peptide, and further provided that when one of R₁ or R₂ is COE₁, the other must be H, or a pharmaceutically acceptable salt thereof.

Preferably, the therapeutic peptide has the formula wherein $A^0 = Gly$, D-Phe, or is deleted; $A^1 = p-Glu$, D-Phe, D-Ala, D-B-Nal, D-Cpa, or D-Asn; $A^2 = Gln$, His, 1-methyl-His, or 3-methyl-His; $A^4 = Ala$;

 $A^5 = Val;$ $A^6 = Sar, Gly, D-Phe, or D-Ala;$ $A^7 = His;$

provided that where R_3 is CH_2-CH_2 , Z_1 is the identifying group of Leu or Phe; or where R_3 is CH_2 , Z_1 is the identifying group of β -Leu or Leu; or where R_3 is $CHOH-CH_2$, Z_1 is the identifying group of Leu or is isopropyl, cyclohexylmethyl, β -naphthylmethyl, or phenylmethyl; provided that V is R_5

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and each R_5 and R_6 is H.

Preferably, the peptide is of the generic formula wherein V is $\rm NHR_6$ where $\rm R_6$ is $\rm NH_2$

Most preferably the therapeutic peptide has the following amino acid formulas:

pGlu-Gln-Trp-Ala-Val-Gly-His-statine-amide; and D-p-Cl-Phe-Gln-Trp-Ala-Val-Gly-His- β -Leu-NH $_2$.

The invention also includes a therapeutic peptide comprising between eight and ten amino acid residues, inclusive, the peptide being an analog of one of the following naturally occurring peptides terminating at the carboxy-terminus with a Met residue:

(a) litorin; (b) neuromedin; (c) the ten amino acid carboxy-terminal region of mammalian gastrin releasing peptide; and (d) the ten amino acid carboxy-terminal region of amphibian bombesin, the therapeutic peptide being of the formula:

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$$R_1$$
 $A^0 - A^1 - A^2 - \text{Trp} - A^4 - A^5 - A^6 - A^7 - W$ R_2

wherein

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- A⁰ = Gly, Nle, α-aminobutyric acid, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β-Nal, or is deleted;
- A¹ = the D- or L-isomer of any of pGlu, Nle, or α -aminobutyric acid, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β -Nal, or is deleted;
- A² = pGlu, Gly, Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, ß-Nal, His, 1-methyl-His, or 3-methyl-His;
- A⁵ = Gln, Asn, Gly, Ala, Leu, Ile, Nle,
 α-aminobutyric acid, Met, Val, p-X-Phe (where
 X = F, Cl, Br, OH, H or CH₃), Trp, Thr, or
 β-Nal;
 - A⁶ = Sar, Gly, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, · Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β-Nal;
 - A' = 1-methyl-His, 3-methyl-His, or His; provided that if A^0 is present, A^1 cannot be pGlu; and if A^0 or A^1 is present, A^2 cannot be pGlu; and when A^0 is deleted and A^1 is pGlu, R_1 must be H and R_2 must be the portion of Glu that forms the imine ring in pGlu; and further provided that W can be:

wherein R_4 is CH_2 -NH, CH_2 -S, CO- CH_2 , or $\mathrm{CH_2-CH_2}$, and each of $\mathrm{Z_1}$ and $\mathrm{Z_2}$, independently, is the identifying group of any one of the amino acids Gly, Ala, Val, Leu, Ile, Ser, Asp, Asn, Glu, Gln, B-Nal, p-X-Phe (where X = H, F, Cl, Br, NO₂, OH or CH₃), Trp, Cys, Met, Pro, HyPro, cyclohexyl-Ala, or cyclohexylmethyl; provided that where R_4 is CH_2-NH and Z_2 is the identifying group of any one of the 10 amino acids Gly, Ala, Val, Leu, Ile, Ser, Asp, Asn, Glu, Gln, p-X-Phe (where X = H, F, Cl, Br, NO₂, OH or CH3), Trp, Cys, Met, Pro, HyPro, or cyclohexylmethyl, \mathbf{Z}_{1} can only be the identifying group of any one of the amino acids Ser, Asp, Glu, Cys, Pro, HyPro, or 15 cylcohexylmethyl; and provided that where R_4 is $ext{CH}_2 ext{-NH}$ and $ext{Z}_1$ is the identifying group of any one of the amino acids Gly, Ala, Val, Leu, Ile, Ser, Asp, Asn, Glu, Gln, p-X-Phe (where X = H, F, Cl, Br, NO_2 , OH or CH_3), Trp, Cys, Met, Pro, or HyPro, Z_2 can only be 20 the identifying group of any one of the amino acids Ser, Asp, Glu, Cys, Pro, HyPro, or cylcohexylmethyl; and V is either OR₅ or

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where each R_8 , R_5 , R_6 , and R_7 , independently, is H, lower alkyl, lower phenylalkyl, or lower naphthylalkyl; and provided that any asymmetric carbon atom can be R, S or a racemic mixture; and further provided that each R_1 and R_2 , independently, is H, C_{1-12} alkyl, C_{7-10} phenylalkyl, COE_1 (where E_1

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is C_{1-20} alkyl, C_{3-20} alkenyl, C_{3-20} alkinyl, phenyl, naphthyl, or C_{7-10} phenylalkyl), or lower acyl, and R_1 and R_2 are bonded to the N-terminal amino acid of the peptide, and further provided that when one of R_1 or R_2 is COE_1 , the other must be H, or a pharmaceutically acceptable salt thereof.

Preferably, the therapeutic peptide has the formula wherein

 $A^0 = Gly$, D-Phe, or is deleted;

10 $A^1 = p-Glu$, D-Phe, D-Ala, D-B-Nal, D-Cpa, or D-Asn;

 $A^2 = Gln$, His, 1-methyl-His, or 3-methyl-His;

 $A^4 = Ala;$

 $A^5 = Val;$

 $A^6 = Sar, Gly, D-Phe, or D-Ala;$

15 $A^7 = His$:

where R_4 is CH_2 -NH, each Z_1 is cyclohexylmethyl or is the identifying group of Leu or Phe; or Z_2 is the identifying group of Met, Leu or Phe.

Most preferably, the therapeutic peptide includes D-B-Nal at position ${\bf A}^1$, where each of ${\bf Z}_1$ and ${\bf Z}_2$, independently, is Leu or Phe.

Examples of such peptides are:

D-G-Nal-Gln-Trp-Ala-Val-Gly-His-Leu ψ [CH2NH]Leu-NH2, or

- D- Ω -Nal-Gln-Trp-Ala-Val-Gly-His-Leu ψ [CH $_2$ NH]Phe-NH $_2$.

 In additon, the therapeutic peptide may include where R $_4$ is CH $_2$ -NH, and the carbon atom bonded to Z $_2$ is of the R configuration. An example of such a peptide is
- D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu ψ [CH₂NH]-D-Phe-NH₂. In addition, the peptide may include where A⁰ or A¹ is a D amino acid, V is OR₄; for example, a methylester derivative such as the therapeutic peptide D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu-Met-methylester.

The invention also includes a therapeutic peptide comprising between seven and nine amino acid residues, inclusive, the peptide being an analog of one of the following naturally occurring peptides terminating at the carboxy-terminus with a Met residue:

(a) litorin; (b) neuromedin; (c) the ten amino acid carboxy-terminal region of mammalian gastrin releasing peptide; and (d) the ten amino acid carboxy-terminal region of amphibian bombesin, the therapeutic peptide being of the formula:

$$A^{0}-A^{1}-A^{2}-Trp-A^{4}-A^{5}-A^{6}-A^{7}-W$$

wherein

- 15 $A^0 = Gly$, Nle, α -aminobutyric acid, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β -Nal, or is deleted;
- the D- or L-isomer of any of pGlu, Nle, or α-aminobutyric acid, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β-Nal, or is deleted;
- pGlu, Gly, Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, β -Nal, His, 1-methyl-His, or 3-methyl-His;
- A^4 = Ala, Val, Gln, Asn, Gly, Leu, Ile, Nle, α -aminobutyric acid, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or β -Nal;

- 14 -

 $A^5 = Gln$, Asn, Gly, Ala, Leu, Ile, Nle, α -aminobutyric acid, Met, Val, p-X-Phe (where X = F, Cl, Br, OH, H or CH₃), Trp, Thr, or β -Nal;

Sar, Gly, or the D-iscmer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or ß-Nal;

 $A^7 = 1$ -methyl-His, 3-methyl-His, or His; provided that if A^0 is present, A^1 cannot be pGlu; and if A^0 or A^1 is present, A^2 cannot be pGlu; and when A^0 is deleted and A^1 is pGlu, R_1 must be H and R_2 must be the portion of Glu that forms the imine ring in pGlu; and further provided that W can be:

wherein \mathbf{Z}_1 is the identifying group of any one of the amino acids Gly, Ala, Val, Leu, Ile, Ser, Asp, Asn, Glu, Gln, p-X-Phe (where X = H, F, Cl, Br, NO_2 , OH or 20 CH_3), Trp , Cys , Met , Pro , or HyPro ; and each R_9 , R_{10} , and R_{11} , independently, is H, lower alkyl, lower phenylalkyl, or lower naphthylalkyl; and provided that any asymmetric carbon atom can be R, S or a racemic mixture; and further provided that each R_1 and R_2 , independently, is H, C_{1-12} alkyl, C_{7-10} phenylalkyl, COE_1 (where E_1 is C_{1-20} alkyl, C_{3-20} alkenyl, C_{3-20} alkinyl, phenyl, naphthyl, or C_{7-10} phenylalkyl), or lower acyl, and R_1 and R_2 are 30 bonded to the N-terminal amino acid of the peptide, and further provided that when one of R_1 or R_2 is \mathtt{COE}_1 , the other must be H, or a pharmaceutically acceptable salt thereof.

WO 90/03980

Preferably, the peptide includes wherein

A⁰ = Gly, D-Phe, or is deleted;

A¹ = p-Glu, D-Phe, D-Ala, D-B-Nal, D-Cpa, or D-Asn;

A² = Gln, His, 1-methyl-His, or 3-methyl-His;

A⁴ = Ala;

A⁵ = Val;

A⁶ = Sar, Gly, D-Phe, or D-Ala;

A⁷ = His;

provided that Z_1 is the identifying group of any one of the amino acids Leu or D or L p-X-Phe (where X = H, F, Cl, Br, NO₂, OH or CH₃); and each R_9 , R_{10} and R_{11} , independently, is H, lower alkyl, lower phenylalkyl, or lower naphthylalkyl.

Most preferably, the peptide includes wherein 15 $^{Z}_{1}$ is Leu, $^{R}_{9}$ is H, and each $^{R}_{10}$ and $^{R}_{11}$ is lower alkyl. Examples of such a peptide are:

 $\label{eq:decomposition} \begin{array}{lll} {\tt D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu-ethylamide;} & or \\ {\tt D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu-NH}_2 \,. \end{array}$

The inventional also includes a therapeutic peptide comprising between six and eight amino acid residues, inclusive, the peptide being an analog of one of the following naturally occurring peptides terminating at the carboxy-terminus with a Met residue:

(a) litorin; (b) neuromedin; (c) the ten amino acid carboxy-terminal region of mammalian gastrin releasing peptide; and (d) the ten amino acid carboxy-terminal region of amphibian bombesin, the therapeutic peptide being of the formula:

wherein

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- Gly, Nle, α -aminobutyric acid, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH_3), Trp, Cys, or $\mathcal{B}\text{-Nal}$, or is deleted;
- $A^1 =$ the D- or L-isomer of any of pGlu, Nle, or α -aminobutyric acid, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, C1, Br, NO_2 , OH, H or CH_3),
- Trp, Cys, or B-Nal, or is deleted; pGlu, Gly, Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO_2 , OH, H or CH_3), Trp, Cys, β -Nal, His, 1-methyl-His, or 3-methyl-His;
- 15 Ala, Val, Gln, Asn, Gly, Leu, Ile, Nle, α -aminobutyric acid, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or ß-Nal;
- $A^5 =$ Gln, Asn, Gly, Ala, Leu, Ile, Nle, 20 α -aminobutyric acid, Met, Val, p-X-Phe (where X = F, Cl, Br, OH, H or CH_3), Trp, Thr, or B-Nal;
 - Sar, Gly, or the D-isomer of any of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, OH, H or CH₃), Trp, Cys, or B-Nal;
- 1-methyl-His, 3-methyl-His, or His; provided that if A^0 is present, A^1 cannot be pGlu; and if A^0 or A^1 is present, A^2 cannot be pGlu; and when ${\tt A}^0$ is deleted and ${\tt A}^1$ is pGlu, ${\tt R}_{\tt l}$ must be H and R_2 must be the portion of Glu that forms the imine ring in pGlu; and further provided that W can be:



wherein each R_{12} and R_{13} , independently, is H, lower alkyl, lower phenylalkyl, or lower naphthylalkyl; provided that any asymmetric carbon atom can be R, S or a racemic mixture; and further provided that each R_1 and R_2 , independently, is H, C_{1-12} alkyl, C_{7-10} phenylalkyl, COE_1 (where E_1 is C_{1-20} alkyl,

 10 $^{\text{C}}_{3-20}$ alkenyl, $^{\text{C}}_{3-20}$ alkinyl, phenyl, naphthyl, or $^{\text{C}}_{7-10}$ phenylalkyl), or lower acyl, and $^{\text{R}}_{1}$ and $^{\text{R}}_{2}$ are bonded to the N-terminal amino acid of the peptide, and further provided that when one of $^{\text{R}}_{1}$ or $^{\text{R}}_{2}$ is $^{\text{COE}}_{1}$, the other must be H, or a pharmaceutically acceptable salt thereof.

Preferably, the therapeutic peptide includes wherein

 $A^0 = Gly$, D-Phe, or is deleted;

 $A^{1} = p-Glu$, D-Phe, D-Ala, D- β -Nal, D-Cpa, or D-Asn;

 $A^2 = Gln$, His, 1-methyl-His, or 3-methyl-His;

 $A^4 = Ala;$

 $A^5 = Val;$

 $A_{-}^{6} = Sar, Gly, D-Phe, or D-Ala;$

 $A^7 = His;$

where each R₁₂ and R₁₃, is H; and each R₁ and R₂, independently, is H, lower alkyl, or lower acyl.

Preferably, where either of N₁₂ or N₁₃ is other than H, A⁷ is His, A⁶ is Gly, A⁵ is Val, A⁴ is Ala, and A² is His; and where either of R₁

30 or R_2 is other than H, A^1 must not be deleted.

The invention also includes a bombesin therapeutic peptide of the formula: pGlu-Gln-Arg-Leu-Gly-Asn-Gln-Trp-Ala-Val-Gly-His-Statine.

5 The invention also includes an effective bombesin antagonistic peptide containing the amino acid formula:

wherein

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A¹ = pGlu, D or L, or is deleted;

15 A² = Gln, Asn, Gly, Ala, Leu, Ile, Nle,
α-aminobutyric acid, Met, Val, Phe, p-X-Phe
(X = F, Cl, Br, OH or CH₃), Trp,
β-naphthylalanine or is deleted;

A³ = Arg, D-Arg, Lys, D-Lys or is deleted;
20 A⁴ = Gln, Asn, Gly, Ala, Leu, Ile, Nle,
α-aminobutyric acid, Met, Val, Phe, p-X-Phe
(X = F, Cl, Br, OH or CH₃), Trp,

ß-naphthylalanine or is deleted; $A^5 = \text{Gln, Asn, Gly, Ala, Leu, Ile, Nle,}$

25 α-aminobutyric acid, Met, Val, Phe, D-Phe, p-X-Phe (X = F, Cl, Br, OH or CH₃), Trp, β-naphthylalanine, D-Ala or is deleted;

A⁶ = Gln, Asn, Gly, Ala, D-Ala, N-Ac-D-Ala, Leu,

Ile, Nle, α-aminobutyric acid, Met, Val, Phe,
p-X-Phe (X = F, Cl, Br, OH or CH₃), Trp,
p-Glu, β-naphthylalanine or is deleted;

- 19 -

A⁷ = Gln, Asn, Gly, Ala, Leu, Ile, Nle, α-aminobutyric acid, Met, Val, Phe, D-Phe, p-X-Phe (X = F, Cl, Br, OH or CH₃), Trp, Lys, His, or β-naphthylalanine;

 $5 A^8 = Trp or Met;$

A⁹ = Gln, Asn, Gly, Ala, Leu, Ile, Nle, α-aminobutyric acid, Met, Val, Phe, p-X-Phe (X = F, Cl, Br, OH or CH₃), Trp, or β-naphthylalanine, D or L;

10 A¹⁰ = Gln, Asn, Gly, Ala, Leu, Ile, Nle, α-aminobutyric acid, Met, Val, Phe, p-X-Phe (X = F, Cl, Br, OH or CH₃), Trp, Thr, or β-naphthylalanine;

 $A^{11} = Gly$, Phe, D or L;

15 $A^{12} = His$, Phe, or p-X-Phe (X = F, Cl, Br, OH, CH₃), D or L;

A¹³ = Gln, Asn, Gly, Ala, Leu, Ile, Nle, α-aminobutyric acid, Met, Val, Phe, p-X-Phe (X = F, Cl, Br, OH or CH₃), Trp, or

ß-naphthylalanine;

A¹⁴ = Gln, Asn, Gly, Ala, Leu, Ile, Nle,
α-aminobutyric acid, Met, Val, Phe, p-X-Phe
(X = F, Cl, Br, OH or CH₃), Trp, or
β-naphthylalanine;

25 provided that

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each R_1 , R_2 , R_3 , and R_4 , independently, is H, C_{1-12} alkyl, C_{7-10} phenylalkyl, COE_1 (where E_1 is C_{1-20} alkyl, C_{3-20} alkenyl, C_{3-20} alkinyl, phenyl, naphthyl, or C_{7-10} phenylalkyl), or $COOE_2$ (where E_2 is C_{1-10} alkyl or C_{7-10} phenylalkyl), and R_1 and R_2 are bonded to the N-terminal amino acid of the peptide, which can be A^1 , A^2 , A^3 , A^4 , A^5 , A^6 , or A^7 , and further provided that when one of R_1 or R_2 is COE_1 or $COOE_2$, the other

must be H, and when one of R_3 or R_4 is COE_1 or $COOE_2$, the other must be H, and further provided that when A^1 = pGlu, R_1 must be H and R_2 must be the portion of Glu that forms the imine ring in pGlu; and for each of the residues A^7 , A^8 , A^9 , A^{11} , A^{12} , and A^{13} , independently, the carbon atom participating in the amide bond between that residue and the nitrogen atom of the alpha amino group of the adjacent amino acid residue may be a carbonyl carbon or may be reduced to a methylene carbon, provided that at least one such carbon atom must be reduced to a methylene carbon,

where the peptide further comprises

 $A^5 = Cys;$

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 A^6 = Cys or a D-isomer of any of the amino acids

described for this position;

 $A^7 = pGlu$, Cys, 1-methyl-His, or 3-methyl-His;

 $A^9 = Cys;$

 $A^{11} = Sar$, or the D-isomer of any of Ala, Val, Gln,

Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂,

20 OH, or CH₃), Trp, Cys, or β-Nal;

 $A^{12} = 1$ -methyl-His, or 3-methyl-His;

and where A¹⁴ may be deleted.

Most preferably, this therapeutic peptide is of the formula:

25 D-p-Cl-Phe-Gln-Trp-Ala-Val-Gly-His-Leuψ[CH₂NH]Phe-NH₂

Preferably, the amino acid sequence of the therapeutic peptides of the formulas described herein are at least 25% homologous with the amino acid sequence of the naturally occurring peptide; most preferably, this homology is at least 50%.

30 this homology is at least 50%.

(Non-peptide bonds in which the peptide bond is reduced are symbolized herein by " $\psi\text{[CH}_2\text{NH]}$ " or " $\psi\text{".}$)

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Antagonists of the invention are useful for treating diseases involving the malignant or benign proliferation of tissue, such as all forms of cancer where bombesin-related or GRP-related substances act as autocrine or paracrine mitotic factors, e.g., cancers of the gastrointestinal tract, pancreatic cancer, colon cancer, lung cancer, particularly the small cell subtype, or breast cancer; or for treating artherosclerosis, and disorders of gastrointestinal tissues related to gastric and pancreatic secretions and motility; for example, for causing the suppression of amylase secretion, or for appetite control.

In the generic formulae given above, when any one of R_1 - R_{13} or R_{15} is an aromatic, lipophilic group, the <u>in vivo</u> activity can be long lasting, and delivery of the compounds of the invention to the target tissue can be facilitated.

The identifying group of an α -amino acid is the atom or group of atoms, other than the α -carbonyl carbon atom, the α -amino nitrogen atom, or the H atom, bound to the asymmetric α -carbon atom. To illustrate by examples, the identifying group of alanine is CH₃, the identifying group of valine is $(CH_3)_2CH$, the identifying group of lysine is $H_3N^+(CH_2)_4$ and the identifying group of phenylalanine is $(C_6H_6)CH_2$. The identifying group of a β - or γ -amino acid is the analagous atom or group of atoms bound to respectively, the β - or the γ -carbon atom. Where the identifying group of an amino acid is not specified it may be α , β , or γ .

Other features and advantages of the invention will be apparent from the following description of the preferred embodiments thereof, and from the claims.

Description of the Preferred Embodiments
We first briefly describe the drawings.

Drawings

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Fig. 1 is a graph of tumor growth curves for NCI-H69 xenografts.

Fig. 2 is a series of amino acid sequences of naturally occurring peptides of which peptides of the invention are analogs.

Fig. 3 is a graph showing the effect of injection of the bombesin analog D-Phe⁶BN(6-13)methylester on bombesin-stimulated pancreatic amylase assay.

We now describe the structure, synthesis, and use of the preferred embodiments of the invention.

15 Structure

The peptides of the invention all have a non-peptide bond in at least one of the indicated positions, except for the statine substituted analogs and ß-leu, such as sta⁸-desLeu⁸-Met⁹ litorin. By non-peptide bond is meant that the carbon atom participating in the bond between two residues is reduced from a carbonyl carbon to a methylene carbon. The peptide bond reduction method which yields this non-peptide bond is described in Coy et al., U.S. patent application, Serial No. 879,348, assigned to the same assignee as the present application, hereby incorporated by reference. Any one of the amino acids in positions 0, 1, 2, and 9 of the litorin antagonists may be deleted from the peptides, and the peptides are still active as antagonists.

The peptides of the invention can be provided in the form of pharmaceutically acceptable salts. Examples of preferred salts are those with therapeutically acceptable organic acids, e.g., acetic,

lactic, maleic, citric, malic, ascorbic, succinic, benzoic, salicylic, methanesulfonic, toluenesulfonic, or pamoic acid, as well as polymeric acids such as tannic acid or carboxymethyl cellulose, and salts with inorganic acids such as the hydrohalic acids, e.g., hydrochloric acid, sulfuric acid, or phosphoric acid. Synthesis of Litorin Analogs

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The synthesis of the litorin antagonist pGlu-Gln-Trp-Ala-Val-Gly-His-Leu ψ [CH2NH]Leu-NH2 follows. Other antagonists of bombesin, litorin, neuromedin, or GRP can be prepared by making appropriate modifications of the following synthetic method.

The first step is the preparation of the intermediate

pGlu-Gln-Trp-Ala-Val-Gly-His(benzyloxycarbonyl)-Leu ψ [CH 2NH]Leu-benzhydrylamine resin, as follows.

Benzhydrylamine-polystyrene resin (Vega Biochemicals, Inc.) (0.97 g, 0.5 mmole) in the chloride ion form is placed in the reaction vessel of a Beckman 990B peptide synthesizer programmed to perform the following reaction cycle: (a) methylene chloride; (b) 33% trifluoroacetic acid (TFA) in methylene chloride (2 times for 1 and 25 min. each); (c) methylene chloride; (d) ethanol; (e) methylene chloride; and (f) 10% triethylamine in chloroform.

The neutralized resin is stirred with alpha-t-butoxycarbonyl(Boc)-leucine and diisopropylcarbodiimide (1.5 mmole each) in methylene chloride for 1 hour, and the resulting amino acid resin is then cycled through steps (a) to (f) in the above wash program. Boc-leucine aldehyde (1.25 mmoles), prepared by the method of Fehrentz and Castro, Synthesis, p. 676 (1983), is dissolved in 5 ml of dry dimethylformamide (DMF) and added to the resin TFA salt

suspension followed by the addition of 100 mg (2 mmoles) of sodium cyanoborohydride (Sasaki and Coy, Peptides 8:119-121 (1987); Coy et al., id.). After stirring for 1 hour, the resin mixture is found to be negative to ninhydrin reaction (1 min.), indicating complete derivatization of the free amino group.

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The following amino acids (1.5 mmole) are then coupled successively in the presence diisopropylcarbodiimide (1.5 mmole), and the resulting amino acid resin is cycled through washing/deblocking steps (a) to (f) in the same procedure as above:

Boc-His(benzyloxycarbonyl), Boc-Gly, Boc-Val, Boc-Ala, Boc-Trp, Boc-Gln (coupled as a 6 M excess of the p-nitrophenylester), and pGlu. The completed resin is then washed with methanol and air dried.

The resin described above (1.6 g, 0.5 mmole) is mixed with anisole (5 ml) and anhydrous hydrogen fluoride (35 ml) at 0°C and stirred for 45 min. Excess hydrogen fluoride is evaporated rapidly under a stream of dry nitrogen, and free peptide is precipitated and washed with ether. The crude peptide is dissolved in a minimum volume of 2 M acetic acid and eluted on a column (2.5 x 100 mm) of Sephadex G-25 (Pharmacia Fine Chemicals, Inc.). Fractions containing a major component by uv absorption and thin layer chromatography (TLC) are then pooled, evaporated to a small volume and applied to a column (2.5 x 50 cm) of octadecylsilane-silica (Whatman LRP-1, 15-20 µm mesh size).

The peptide is eluted with a linear gradient of 0-30% acetonitrile in 0.1% trifluoroacetic acid in water. Fractions are examined by TLC and analytical high performance liquid chromatography (HPLC) and pooled to give maximum purity. Repeated lyophilization of the

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solution from water gives 60 mg of the product as a white, fluffy powder.

The product is found to be homogeneous by HPLC and TLC. Amino acid analysis of an acid hydrolysate confirms the composition of the peptide. The presence of the Leu ψ [CH₂-NH]Leu bond is demonstrated by fast atom bombardment mass spectrometry. pGlu-Gln-Trp-Ala-Val-Gly-His-Phe ψ [CH₂NH]Leu-NH₂ and

pGlu-Gln-Trp-Ala-Val-Gly-His-Leu ψ [CH2NH]Leu-NH2 or other peptides are prepared in similar yields in an analogous fashion by appropriately modifying the above procedure.

Solid phase synthesis of D-Phe¹,

Leu⁸ \(\text{CH}_2\text{NH} \] D-Phe⁹-litorin, D-Phe-Gln-Trp-Ala-Val-Gly- His-Leu\(\text{CH}_2\text{NH} \] -D-Phe-NH₂, was carried out as follows:

Boc-D-Phe-Gln-Trp-Ala-Val-Gly-His(tosyl)-Leu\(\text{CH}_2\text{NH} \] -D-Phe-benzhydrylamine resin was synthesized first.

Benzhydrylamine-polystyrene resin (Advanced ChemTech, Inc.) (1.25 g, 0.5 mmole) in the chloride ion form is placed in the reaction vessel of an Advanced ChemTech ACT 200 peptide synthesizer programmed to perform the reaction cycle described as steps (a) through (f) above.

The neutralized resin is stirred with

Boc-D-phenylalanine and diisopropylcarbodiimide (1.5 mmole
each) in methylene chloride for 1 h and the resulting
amino acid resin is then cycled through steps (a) to (g)
in the above wash program. The Boc group is then removed
by TFA treatment. Boc-leucine aldehyde (1.25 mmoles),
prepared by the method of Fehrentz and Castro (1), is
dissolved in 5 ml of dry DMF and added to the resin TFA
salt suspension followed by the addition of 100 mg (2

mmoles) of sodium cyanoborohydride (2,3). After stirring for 1 h, the resin mixure is found to be negative to ninhydrin reaction (1 min) indicating complete derivatization of the free amino group.

The following amino acids (1.5 mmole) are then coupled successively by the same procedure: Boc-His(benzyloxycarbonyl), Boc-Gly, Boc-Val, Boc-Ala, Boc-Trp, Boc-Gln (coupled in the presence of 1 equiv. hydroxybenzotriazole), Boc-D-Phe (coupled in the presence 10 of 1 equiv. hydroxybenzotriazole). After drying, the peptide resin weighed 1.93 g.

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The resin (1.93 g, 0.5 mmole) is mixed with anisole (5 ml) and anhydrous hydrogen fluoride (35 ml) at 0°C and stirred for 45 min. Excess hydrogen fluoride is evaporated rapidly under a stream of dry nitrogen and free peptide precipitated and washed with ether. The crude peptide is dissolved in a minimum volume of 2 M acetic acid and eluted on a column (2.5 x 100 mm) of Sephadex G-25. Fractions containing a major component by uv absorption and thin layer chromatography are then pooled, 20 evaporated to a small volume and applied to a column (2.5 \times 50 cm) of Vydac octadecylsilane (10-15 uM). This is eluted with a linear gradient of 15-45% acetonitrile in 0.1% trifluoroacetic acid in 25 water. Fractions are examined by thin layer chromatography and analytical high performance liquid chromatography and pooled to give maximum purity. Repeated lyophlization of the solution from water gives 120 mg of the product as a white, fluffy powder.

Other peptides, e.g., D-p-Cl-Phe-Gln-Trp-Ala-Val-Gly-His-Leuw[CH2NH]-D-Phe-NH2, may be synthesized using essential the same procedure.

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Synthesis of Leu⁸ w[CH₂NH] D-Phe⁹ Litorin

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Solid phase synthesis of

D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu ψ [CH2NH]-D-Phe-NH2 was carried out as follows:

Boc-D-Phe-Gln-Trp-Ala-Val-Gly-His(tosyl)-Leuw[CH2NH]-D-Phe-benzhydrylamine resin was synthesized first.

Benzhydrylamine-polystyrene resin (Advanced ChemTech, Inc.) (1.25 g, 0.5 mmole) in the chloride ion form is placed in the reaction vessel of an Advanced ChemTech ACT 200 peptide synthesizer programmed to perform the following reaction cycle: (a) methylene chloride; (b) 33% trifluoroacetic acid in methylene chloride (2 times for 1 and 25 min each); (c) methylene chloride; (d) ethanol; (e) methylene chloride; (f) 10% triethylamine in chloroform.

The neutralized resin is stirred with Boc-D-phenylalanine and diisopropylcarbodiimide (1.5 mmole each) in methylene chloride for 1 h and the resulting amino acid resin is then cycled through steps (a) to (g) in the above wash program. The Boc group is then removed by TFA treatment. Boc-leucine aldehyde (1.25 mmoles), prepared by the method of Fehrentz and Castro (1), is dissolved in 5 ml of dry DMF and added to the resin TFA salt suspension followed by the addition of 100 mg (2 mmoles) of sodium cyanoborohydride (2,3). After stirring for 1 h, the resin mixure is found to be negative to ninhydrin reaction (1 min) indicating complete derivatization of the free amino group.

The following amino acids (1.5 mmole) are then coupled successively by the same procedure:

Boc-His(benzyloxycarbonyl), Boc-Gly, Boc-Val, Boc-Ala, Boc-Trp, Boc-Gln (coupled in the presence of 1 equiv. hydroxybenzotriazole), Boc-D-Phe (coupled in the presence of 1 equiv. hydroxybenzotriazole). After drying, the peptide resin weighed 1.93 g.

The resin (1.93 g, 0.5 mmole) is mixed with anisole (5 ml) and anhydrous hydrogen fluoride (35 ml) at 0°C and stirred for 45 min. Excess hydrogen fluoride is evaporated rapidly under a stream of dry nitrogen and 5 free peptide precipitated and washed with ether. crude peptide is dissolved in a minimum volume of 2 M abetic acid and eluted on a column (2.5 x 100 mm) of Sephadex G-25. Fractions containing a major component by uv absorption and thin layer chromatography are then 10 pooled, evaporated to a small volume and applied to a column (2.5 \times 50 cm) of Vydac octadecylsilane (10-15 uM). This is eluted with a linear gradient of 15-45% acetonitrile in 0.1% trifluoroacetic acid in water. Fractions are examined by thin layer chromatography and analytical high performance liquid 15 chromatography and pooled to give maximum purity. Repeated lyophlization of the solution from water gives 120 mg of the product as a white, fluffy powder.

The product is found to be homogeneous by hplc and tlc. Amino acid analysis of an acid hydrolysate confirms the composition of the octapeptide. The presence of the Leuw[CH2NH] peptide bond is demonstrated by fast atom bombardment mass spectrometry.

25 Synthesis of D-Phe 1-Des-Met 9 Litorin

Solid phase synthesis of D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu-NH $_2$ was carried out as follows.

Step (1): Benzhydrylamine-polystyrene resin

(Advanced ChemTech, Inc. (0.62 gm, 0.25 mmole) in the chloride ion form is placed in the reacton vessel of an ACT 200 peptide synthesizer programmed to perform the following reaction cycle: (a) methylene chloride; (b)

33% trifluoroacetic acid in methylene chloride (2 times

- 29 -

for 1 and 25 min each); (c) methylene chloride; (d) ethanol; (e) methylene chloride; (f) 10% triethylamine in chloroform.

Boc-leucine and diisopropylcarbodiimide (1.5 mmole each) in methylene chloride for 1 hr and the resulting amino acid resin is then cycled through steps (a) to (g) in the above wash program. The following amino acids (1.5 mmole) are then coupled successively by the same procedure: Boc-His (benzyloxycarbonyl, Boc-Gly, Boc-Val, Boc-Ala, Boc-Trp, Boc-Gln (coupled as a 6M excess of the p-nitrophenylester, and pGlu (coupled in the presence of hydroxzybenzotriazole). After drying, the peptide resin

weighed 0.92 q. 15 Step (2): The resin (0.92 g) is then mixed with anisole (5 ml), dithiothreitol (200 mg) and anhydrous hydrogen fluoride (35 ml) at 0° C and stirred for 45 Excess hydrogen fluoride is evaporated rapidly under a stream of dry nitrogen and free peptide precipitated and washed with ether. The crude peptide 20 is dissolved in a minimum volume of 2 M acetic acid and eluted on a column (2.5 \times 100 cm) of Sephadex G-25. Fractions containing a major component by UV absorption and thin layer chromatography are then pooled, 25 evaporated to a small volume and applied to a column (2.5 \times 50 cm) of Vydac octadecylsilane (10-15 microM). The column is eluted with a linear gradient of 0-30% acetonitrile in 0.1% trifluoroacetic acid in water. Fractions are examined by thin layer chromatography and 30 pooled to give maximum purity. Repeated lyophilization of the solution from water gives a white, fluffy powder;

this product is found to be homogeneous by hplc and

confirms the compositon of the peptide.

Amino acid analysis of an acid hydrolysate

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Synthesis of

D-Nal-Gln-Trp-Ala-Val-Gly-His-Leu-NH $_2$ was accomplished using the same procedure as described above (0.62 g, 0.25 mmole of benzyhydrylamine resin in step (1), and 0.92 g in step (2)).

Synthesis of

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N-acetyl-D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu-NH₂ was accomplished using the same procedure as that described above, using 0.62 g (0.25 mmole) of benzhydrylamine resin in step (1), and mixing 0.92 g of the resin with anisole in step(2), except that the final Boc group was removed and the resin acetylated with acetic anhydride in methylene chloride.

Synthesis of Sta⁸-Des-Met⁹ Litorin

15 A statine, AHPPA, or ACHPA residue can be substituted in place of any two amino acids of the peptide, where the peptide contains only peptide bonds. For example, sta⁸-des Met⁹ litorin was prepared in an analagous fashion by first coupling statine to the 20 resin and then proceeding with the addition of Boc-His(benzylocarbonyl). Statine or Boc-statine can be synthesized according to the method of Rich et al., 1978, J. Organic Chem. 43; 3624; and Rich et al., 1980, J. Med. Chem. 23: 27, and AHPPA and ACHPA can be synthesized according to the method of Hui et al., 1987, 25 J. Med. Chem. 30: 1287; Schuda et al., 1988, J. Org. Chem. 53:873; and Rich et al., 1988, J. Org. Chem. 53:869.

Solid-phase synthesis of the peptide BIM-26120,

pGlu-Gln-Trp-Ala-Val-Gly-His-Sta-NH₂, was accomplished through the use of the following procedures in which alpha-t-butoxycarbonyl statine (prepared by the procedure of Rich et al., J. Org. Chem. 1978, 43, 3624) is first coupled to methylbenzhydrylamine-polystyrene

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resin. After acetylation, the intermediate p-Glu-Gln-Gln-Trp-Ala-Val-Gly-His(benzyloxycarbonyl)-Stamethylbenzhydrylamine resin is prepared. The synthetic procedure used for this preparation follows in detail:

1. Incorporation of alpha-t-butoxycarbonyl statine on methylbenzhydrylamine resin.

Methylbenzhydrylamine-polystyrene resin (Vega Biochemicals, Inc.) (1.0 g, 0.73 mmol) in the chloride ion form is placed in the reaction vessel of a Vega 250C Coupler peptide synthesizer. The synthesizer was programmed to perform the following reactions: (a) methylene chloride; (b) 10% triethylamine in chloroform; (c) methylene chloride; and (d) dimethylformamide.

with the preformed active ester made from alpha-t-butoxycarbonyl statine (1.46 mmol), diisopropyl carbodiimide (2 mmol), and hydroxybenzotriazole hydrate (1.46 mmol in dimethylformamide at 0° C. for one hour. The resulting amino acid resin is washed on the synthesizer with dimethylformamide and then methylene chloride. The resin mixture at this point was found by the Kaiser ninhydrin test (5 minutes) to have an 84% level of statine incorporation on the resin.

Acetylation was performed by mixing the amino-acid resin for 15 minutes with N-acetyl imidazole (5 mmol) in methylene chloride. Derivatization to the 94-99% level of the free amino groups of the resin was indicated by the Kaiser ninhydrin test (5 minutes). The Boc-statine-resin is then washed with methylene chloride.

2. Couplings of the Remaining Amino Acids.

The peptide synthesizer is programmed to perform the following reaction cycle: (a) methylene chloride; (b) 33% trifluoroacetic acid (TFA) in methylene chloride (2 times for 5 and 25 min. each); (c)

methylene chloride; (d) isopropyl alcohol; (e) 10% triethylamine in chloroform; and (f) methylene chloride.

The following amino acids (2.19 mmol) are then coupled successively by diisopropyl carbodiimide (4 mmol) alone or diisopropyl carbodiimide (4 mmol) plus hydroxybenzotriazole hydrate (1.47 or 0.73 mmol) and the resulting peptide-resin is washed on the synthesizer with dimethylformamide and then methylene chloride, and then cycled through the washing and deblocking steps (a) to (f) in the procedure described above.

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Boc-His (benzyloxycarbonyl) (coupled in the presence of 2 equivalents hydroxybenzotriazole);
Boc-Gly; Boc-Val; Boc-Ala and Boc-Trp (coupled as the preformed hydroxybenzotriazole active esters made by reaction at 0° C for one hour with 1 equivalent hydroxybenzotriazole hydrate); Boc-Gln and pGlu (also coupled as the preformed active esters of hydroxybenzotriazole made by reaction at 0° C for one hour with 1 equivalent hydroxybenzotriazole hydrate). The completed peptide-resin is then washed with methanol and air dried.

The peptide-resin described above (1.60 g, 0.73 mmol) is mixed with anisole (2.5 mL), dithioerythreitol (50 mg), and anhydrous hydrogen fluoride (30 mL) at 0°C. for one hour. Excess hydrogen fluoride is evaporated rapidly under a stream of dry nitrogen, and the free peptide is precipitated and washed with ether. The crude peptide is dissolved in 100 mL of 1 M acetic acid and the solution is then evaporated under reduced pressure. The crude peptide is dissolved in a minimum volume of methanol/water 1/1 and triturated with 10 volumes of ethyl acetate.

The triturated peptide is applied to a column (9.4 mm I.D. \times 50 cm) of octadecylsilane-silica (Whatman

Partisil 10 ODS-2 M 9). The peptide is eluted with a linear gradient of 20-80% of 20/80 0.1% trifluoroacetic acid/acetonitrile in 0.1% trifluoroacetic acid in water. Fractions are examined by TLC and analytical high performance liquid chromatography (HPLC) and pooled to give maximum purity. Lyophilization of the solution from water gives 77 mg of the product as a white fluffy powder.

Other compounds including D-Cpa¹, ß-leu⁸,

desMet⁹ Litorin can be prepared as above and tested for effectiveness as agonists or antagonists in the following test program.

Phase 1 - 3T3 Peptide Stimulated [3H] Thymidine
Uptake Assay

15 Cell Culture. Stock cultures of Swiss 3T3

cells are grown in Dulbecco's Modified Eagles Medium
(DMEM) supplemented with 10% fetal calf serum in
humidified atmosphere of 10% CO2/90% air at 37°C. For
experimental use, the cells are seeded into 24-well

cluster trays and used four days after the last change
of medium. The cells are arrested in the G1/G0 phase of
the cell cycle by changing to serum-free DMEM 24 hours
prior to the thymidine uptake assay.

25 twice with 1ml aliquots of DMEM (-serum) then incubated with DMEM (-serum), 0.5µM [methyl-3H] thymidine (20Ci/mmole, New England Nuclear); bombesin (3nM), and initially four concentrations of the test compounds (1, 10, 100, 1000nM) in a final volume of 1.0 ml. After 28 hours at 37°C, [methyl-3H] thymidine incorporation into acid-insoluble pools is assayed as follows. The cells are washed twice with ice-cold 0.9% NaCl (1ml aliquots), and acid soluble radioactivity is removed by a 30 min. (4°C) incubation with 5% trichloroacetic acid

(TCA). The cultures are then washed once (Iml) with 95% ethanol and solubilized by a 30 min. incubation (Iml) with 0.1N NaOH. The solubilized material is transferred to vials containing 10ml ScintA (Packard), and the radioactivity is determined by liquid scintillation spectrometry.

Phase 2 - Small Cell Carcinoma (SCLC) - Bombesin

Stimulated [3H] Thymidine Uptake Assay

Cell Culture. Cultures of the human cell

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Assay of DNA Synthesis. Bombesin (lnM),
0.5µM [methyl-3H] thymidine (20 Ci/mmole, New
England Nuclear), and four concentrations of the test
compounds (1, 10, 100, 1000nM) are added to the cultures
to achieve a final volume of 0.5 ml. After a 28 hr
incubation at 37°C, the cells are collected onto GF/B
glass fiber filters, and the DNA is precipitated with
ice-cold TCA. [3H] thymidine incorporation into
acid-insoluble fractions of DNA is determined by liquid
scintillation spectrometry.

Phase 3 - Peptide-Induced Pancreatitis

Male, Sprague-Dawley rats (250g) are used for these experiments. The test compound, or 0.9% NaCl is administered s.c. 15 min. prior to the bombesin injection. Bombesin injections are given s.c. at a dose of 10 μg/kg, and blood samples are obtained at 1 hr.30 min., 3hr. and 6hr. Plasma amylase concentration are determined by the Pantrak Amylase test.

- 35 -

Phase 4- In Vitro Inhibition of [125 I] Gastrin Releasing Peptide (GRP) Binding to Bombesin Receptors

Membranes from various tissues (rat brain, rat pancreas, rat anterior pituitary, SCLC, 3T3 cells) are 5 prepared by homogenization in 50mM TrisHCl containing 0.1% bovine serum albumin and 0.1mg/ml bacitracin followed by two centrifugations (39,000xgx15 min., 4°C) with an intermediate resuspension in fresh buffer. For assay, aliquots (0.8ml) are incubated with 0.5nM 10 $[^{125}I]GRP$ ('2000 Ci/mmol, Amersham Corp.) and various concentrations of the test compounds in a final volume of 0.5ml. After a 30 minute incubation at 4°C, the binding reaction is terminated by rapid filtration through Whatman GF/C filters that have been pre-soaked 15 in 0.3% aqueous polethyleneimine to reduce the level of nonspecific binding. The filters and tubes are washed three times with 4ml aliquots of ice-cold buffer, and the radioactivity trapped on the filters is counted by gamma-spectrometry. Specific binding is defined as the 20 total $[^{125}\text{I}]\text{GRP}$ bound minus that bound in the presence of 1000nM bombesin.

Phase 5- Inhibition of Gastrin Release

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The stomachs of anesthetized rats are perfused with saline collected over 15 minute periods via pyloric cannulation while the test peptide is infused through the femoral vein for periods between 0 and 150 minutes. Phase 6- In Vivo Antitumor Activity

NCI-H69 small cell lung carcinoma cells were transplanted from <u>in vitro</u> culture by implanting each animal with the equivalent of 5 confluent 75 cm² tissue culture flasks in the right flank. <u>In vitro</u> NCI-H69 cells grow as a suspension of cellular aggregates. Therefore, no attempt was made to

disaggregate the cell agglomerates by physical or chemical means. Tumor size was calculated as the average of two diameters, i.e., (length and width/2) mm.

5 Results of Assays of Test Peptides

A number of analogs of bombesin or GRP, each containing a non-peptide bond or a statine, AHPPA, or ACHPA residue, can be synthesized and tested in one or more of the above-described Phase 1 - 6 assays; the results of Phase 1 and 2 tests are given in Table 1 10 attached hereto. Table I shows formulas for the non-peptide analogues and results of in vitro inhibition of [125] GRP binding to 3T3 fibroblast bombesin receptors, and bombesin-stimulated [3H]Thymidine uptake by cultured 3T3 cells. (3T3 GRP receptor and thymidine uptake data are expressed in IC50 (nM).) Table 1 also gives results for non-peptide bond-containing analogs of one other naturally-occurring peptide, Neuromedin C, whose C-terminal seven amino acids are similar to those of bombesin and GRP. 20 Table I, "Litorin" indicates a 9 residue peptide analog or its derivative, whereas "Neuromedin C" indicates a 10 residue analog or its derivative.)

In Table I, the position of the non-peptide

bond is indicated by the position of the symbol

w[CH2NH]; i.e., w[CH2NH] is always shown

preceding the amino acid which, in that peptide, is

bonded to the amino acid N-terminal to it via the

non-peptide bond. Where no amino acid is specified

under "structure", the non-peptide bond links the two

peptides represented by the numbers given as

post-scripts.

In Table I, it can be seen that a preferred placement of the non-peptide bond in litorin analogs is

- 37 -

at the ${\rm A}^8$ - ${\rm A}^9$ position; two of the most active analogs (as indicated by a low GRP receptor IC50 value) are BIM-26100 (Phe $^8\psi[{\rm CH_2NH}]{\rm Leu}^9$) and BIM-26101 (Leu $^8\psi[{\rm CH_2NH}]{\rm Leu}^9$).

In addition, as shown in Table I, BIM-26113 (D-Phe 1 , Leu $^8\psi$ [CH $_2$ NH]Leu 9) and BIM-26114 (D-Nal 1 , Leu $^8\psi$ [CH $_2$ NH]Leu 9) are active in the 3T3 GRP receptor binding and thymidine uptake assays. Most notably, BIM-26136 (D-Nal 1 ,

Leu $^8\psi[\text{CH}_2\text{NH}]\text{Phe}^9)$, which contains amino and carboxy terminal aromatic residues that are capable of forming a hydrophobic interaction, is the most potent analog. Finally, when statine or β -leucine replaces the A^8 and A^9 residues of litorin, the resultant analogs BIM-26120 and BIM-26182 are also potent antagonists.

Table I also shows that Neuromedin C analogs containing a non-peptide bond between residues A^9 - A^{10} , e.g., BIM-26092, 26095, 26106, and 26107, are antagonists when tested in the 3T3 GRP receptor and thymidine uptake assays.

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Table 1 also gives negative results for analogs of Neuromedin C, e.g., BIM-26108. Thus the non-peptide bond placement guidelines given herein should be used in conjunction with the routine assays described above to select useful antagonists.

Bombesin and Bombesin analogs have been shown to inhibit the effect of interleukin-2 (IL-2) (Fink et al., 1988, Klin. Wochenschr. 66, Suppl. 13, 273). Since IL-2 causes T lymphocytes to proliferate, it is possible that litorin antagonists may prevent the inhibitory effect of Bombesin or its analogs on IL-2. IL-2 stimulated lymphocytes are capable of effectively lysing small cell lung carcinoma cells in vitro. Although Bombesin antagonists have a direct antiproliferative

effect on neoplastic tissues, they may also favor proliferation of lymphocytes having lytic activity for small cell lung carcinoma.

These observations prompted us to evaluate the effect of BIM-26100 on the <u>in vivo</u> growth of the SCLC tumor cell line described in Phase 6. Twenty athymic nude females, 5 to 6 weeks of age, were implanted on day 0 with the NCI-H69 human SCLC, individually identified and then randomized into the following vehicle control and test groups:

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Group No. Treatment No. Animals

1 Saline vehicle treated control:

0.2 ml, s.c. inf., b.i.d., QD1-28 10

2 BIM-26100:

50ug/inj., s.c., b.i.d., QD1-28 5

3 BIM-26100:

50ug/inj., s.c. inf., b.i.d., QD1-28 5

(s.c. = subcutaneously; inj. = injected; b.i.d. = twice per day; QD1-28 = daily treatment, on days 1 - 28.)

Growth of NCI-H69 xenografts and the tumor growth inhibitory activity of the bombesin antagonist BIM-26100

(pGlu-Gln-Trp-Ala-Val-Gly-His-Pheψ[CH₂NH]Leu-NH₂) are illustrated as tumor growth curves in Fig. 1, and relative tumor sizes in Table II. Administration of BIM-26100 as a s.c. infusion around the tumor significantly inhibited tumor growth. The effectiveness of the antitumor activity of BIM-26100 is evident in view of the large inoculum of NCI-H69 tumor cells (i.e., the equivalent of 5 confluent 75 cm² cell culture flasks per animal) and the agglomerated condition of the cells. In confluent flasks, NCI-H69 agglomerates are macroscopically visible and together resemble a metastatic tumor colony. Many such tumor colonies were implanted per animal. The dose of BIM-26100 was

- 39 -

arbitrarily selected on the basis of compound availability and is not optimal. Higher doses of BIM-26100 may be administered, as indicated by body weight gain (minus tumor weight) gain during the course of treatment (Table III). This suggest BIM-26100 completely lacks local or systemic toxicity and is useful therapeutically as an anti-growth factor with anti-tumor effects.

Fig. 3 shows the effect of the bombesin 10 antagonist ${\tt D-p-Cl-Phe-Gln-Trp-Ala-Val-Gly-His-Leu\psi[CH_2NH]Phe-NH_2}$ on bombesin-stimulated amylase secretion in the rat. The results show that this analog is a potent antagonist; 5 nM of the analog can inhibit the secretion of anylase stimulated by 0.5 nM of bombesin for 150 minutes after bolus injection. Use

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The peptides of the invention may be administered to a mammal, particularly a human, in one of the traditional modes (e.g., orally, parenterally, transdermally, or transmucosally), in a sustained release formulation using a biodegradable biocompatible polymer, or by on-site delivery (e.g., in the case of anti-cancer bombesin to the lungs) using micelles, gels and liposomes.

The bombesin antagonists of the invention are suitable for the treatment of all forms of cancer where bombesin-related substances act as autocrine or paracrine mitotic agents, particularly small-cell lung carcinoma. The peptides can also be used for the inhibition of gastric acid secretion and motility disorders of the GI tract, the symptomatic relief and/or treatment of exocrine pancreatic adenocarcinoma, and the restoration of appetite to cachexic patients.

- 40 -

peptides can be administered to a human patient in a dosage of $0.5~\mu g/kg/day$ to 5~mg/kg/day. For some forms of cancer, e.g., small cell lung carcinoma, the preferred dosage for curative treatment is 250mg/patient/day.

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- 41 -

Table 1

	- CO T		
		3T3 GRP	Thym.
		Receptor	Uptake
<u>Code</u>	Structure	IC50(nM)	IC50(nM)
BIM-26092	Gly-Asn-His-Trp-Ala-Val-Gly-		
	His-Leuw[CH2NH]Leu-NH2		
	Neuromedin C	242	466
	•	2.12	±00
BIM-26095	pGlu-Gln-Trp-Ala-Val-D-Ala-		
	His-Leuw[CH2NH]Leu-NH2		
• -	Litorin	2623	1209
BIM-26100	pGlu-Gln-Trp-Ala-Val-Gly-His-		•
	Phew[CH2NH]Leu-NH2		
	Litorin	23	26
		23	26
BIM-26101	pGlu-Gln-Trp-Ala-Val-Gly-His-		
	Leuw[CH2NH]Leu-NH2		
	Litorin	118	296
BIM-26105	D-Ala-Asn-His-Trp-Ala-Val-		-
	D-ALa-His-Leuw[CH2CH]Leu-NH2	•	,
	Neuromedin C	107	107
			20,
BIM-26106	desGly-D-Ala-His-Trp-Ala-Val-		
	D-Ala-His-Leuw[CH2NH]Met-NH2		
	Neuromedin C	401	354
		401	334
BIM-26107	D-Phe-His-Trp-Ala-Val-Gly-		
	His-Leuψ[CH ₂ NH]Leu-NH ₂		-
	Neuromedin C	199	154
	-	エフフ	154

_ 42 _

Table 1 (cont'd)

	Table I (Court a)		
		3T3 GRP	Thym.
		Receptor	Uptake
Code	Structure	IC50(nM)	IC50(nM)
BIM-26108	N-Ac-D-Ala-His-Trp-Ala-Val-		
	Gly-His-Leuw[CH2NH]Leu-NH2		
	Neuromedin C	841	>1000
BIM-26113	D-Phe-Gln-Trp-Ala-Val-Gly-		
	His-Leuψ[CH ₂ NH]Leu-NH ₂		
	Litorin	5.8	9
BIM-26114	D-Nal-Gln-Trp-Ala-Val-Gly-		
	His-Leuψ[CH ₂ NH]Leu-NH ₂		
	Litorin	23.5	28
BIM-26120	pGlu-Gln-Trp-Ala-Val-Gly-		
	His-Sta-NH ₂		
	Litorin	150	165
BIM-26122	D-Phe-Gln-Trp-Ala-Val-Gly-		
	His-Leu-NH ₂		
	Litorin	5.9	28.6
		3.3	20.0
BIM-26136	D-Nal-Gln-Trp-Ala-Val-Gly-His-		
	Leuw[CH2NH]Phe-NH2		
	Litorin	1.4	3.3
	·		
BIM-26182	D-Cpa-Gln-Trp-Ala-Val-Gly-His-	0.88	4.77
	B-Leu-NH ₂ Litorin		

Table II

IN YIYO, TUMOR INHIBITORY ACTIVITY OF THE BOMBESIN ANTAGONIST BIM-26100; NCL-1162 HUMAN SCLC

1 Vehicle treated control, 0.2ml, s.c. inf. b.i.d., QD1-28 2 BIM-26100, S0µg/mj. s.c., b.i.d., QD1-28 3 BIM-26100, S0µg/mj. s.c. inf., b.i.d., QD1-28 7.6±1.56** 70 13.7±0.67* 86	Group No. Treatment	Tumor Size ¹ Day 18 (mm)	% Test/Control	Tumor Size Day 28 (mm)	% Test/Control
BIM-26100, 50µg/mj. ş.c., b.i.d., QD1-28 10.1±1.47 93 17.3±1.96 BIM-26100, 50µg/mj. s.c. inf., b.i.d., QD1-28 7.6±1.56** 70 13.7±0.67*		10.9±1.82		15.942.27	
inf., b.i.d., QD1-28 7.6±1.56** 70 13.7±0.67*	BIM-26100, 50pg/inf., s.c	10.1±1.47	93	17.3±1.96	108
		7.6±1.56**	70	13.7±0.67	98

¹Data reported as means ± SD on 10 animals in the control and 5 in test groups. Student's t Test significance of difference from control: *p<0.05; **p<0.01.

Table III

EFFECT OF TUMOR GROWTH AND BIM-26100 TREATMENT ON BODY WEIGHT:
LACK OF SYSTEMIC TOXICITY

No. Treatment Day 0 Weight(gm) Body Weight(gm) 1 Vchicle treated control, 0.2ml, s.c. inf., 17.3 19.6 19.7 2 BIM-26100, 50µg/inj., s.c., b.i.d., QD1-28 16.9 19.2 19.1 3 BIM-26100, 50µg/inj., s.c. inf., b.i.d., QD1-28 17.7 20.4 21.1	Gro	do	Rody Weightfamil	,	
st. 0.2ml, s.c. inf., s.c., b.i.d., QD1-28 s.c. inf., b.i.d., QD1-28	g		Day 0	body Weight (gm) Day 18	Body Weight(gm)
s.c., b.i.d., QDI-28 16.9 19.2 s.c. inf., b.i.d., QDI-28 17.7 20.4	-				797.70
s.c., b.i.d., QDI-28 16.9 19.2 s.c. inf., b.i.d., QDI-28 17.7 20.4			17.3	19.6	19.7
s.c. inf., b.i.d., QD1-28 17.7 20.4	8	-	16.9	. 19.2	19.1
	ന	•	17.7	20.4	21.1

Body weights are presented as means of 10 animals in the control and 5 in test groups.

Tumor weights calculated from 2 largest diameters in mm converted to mgs using the formula for an ellipsoid (length x width 2/2)mgs, were subtracted from total body weights,

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Claims

1 A linear peptide which is an analog of naturally occurring, biologically active amphibian 2 bombesin or mammalian GRP having an active site and a 3 4 binding site responsible for the binding of said peptide 5 to a receptor on a target cell, cleavage of a peptide bond in said active site of naturally occurring bombesin 6 or GRP being unnecessary for in vivo biological 7 8 activity, said analog having one of the following modifications: (a) a deletion of a residue within said 10 active site and a modification of a residue outside of 11 said active site, or (b) a replacement of one or two 12 residues within said active site with a synthetic amino acid, said analog being capable of binding to said 13 receptor so that said analog is capable of acting as a 14 15 competitive inhibitor of said naturally occurring 16 peptide by binding to said receptor and, by virtue of 17 one of said modifications, failing to exhibit the in vivo biological activity of said naturally occurring 18 19 bombesin or GRP.

- 2. The linear peptide of claim 1 wherein said active site comprises at least one amino acid in the carboxy terminal half of the peptide, said linear peptide including said amino acid in its carboxy terminal half.
- 3. The linear peptide of claim 1 wherein said active site includes at least one amino acid in the amino terminal half of the peptide, said linear peptide including said amino acid in its amino terminal half.

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1
                   A therapeutic peptide comprising between
 2
     seven and nine amino acid residues, inclusive, said
     peptide being an analog of one of the following
 3
     naturally occurring peptides terminating at the
 4
     carboxy-terminus with a Met residue: (a) litorin; (b)
 5
     neuromedin; (c) the ten amino acid carboxy-terminal
6
     region of mammalian gastrin releasing peptide; and (d)
7
     the ten amino acid carboxy-terminal region of amphibian
8
     bombesin, said therapeutic peptide being of the formula:
10
             A^{0}-A^{1}-A^{2}-Trp-A^{4}-A^{5}-A^{6}-A^{7}-W
11
12
13
    wherein
    A^0 =
14
              Gly, Nle, \alpha-aminobutyric acid, or the
15
              D-isomer of any of Ala, Val, Gln, Asn, Leu,
16
              Ile, Met, p-X-Phe (where X = F, C1, Br, NO<sub>2</sub>,
17
              OH, H or CH3), Trp, Cys, or B-Nal, or is
18
              deleted:
19
              the D- or L-isomer of any of pGlu, Nle, or
20
              \alpha-aminobutyric acid, or the D-isomer of any
21
              of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe
22
              (where X = F, C1, Br, NO<sub>2</sub>, OH, H or CH<sub>3</sub>),
23
              Trp, Cys, or B-Nal, or is deleted;
24
             pGlu, Gly, Ala, Val, Gln, Asn, Leu, Ile, Met,
25
             p-X-Phe (where X = F, Cl; Br, NO_2, OH, H or
26
             CH_3), Trp, Cys, \beta-Nal, His, 1-methyl-His, or
27
             3-methyl-His;
28
            Ala, Val, Gln, Asn, Gly, Leu, Ile, Nle,
29
             \alpha-aminobutyric acid, Met, p-X-Phe (where X =
```

F, Cl. Br. NO_2 , OH, H or CH_3), Trp. Cys. or

30

31

B-Nal;

WO 90/03980

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32
                Gln, Asn, Gly, Ala, Leu, Ile, Nle,
 33
                \alpha-aminobutyric acid, Met, Val, p-X-Phe (where
 34
                X = F, C1, Br, OH, H or CH_2), Trp, Thr, or
 35
                B-Nal;
 36
                Sar, Gly, or the D-isomer of any of Ala, Val,
 37
                Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F,
 38
                C1, Br, NO_2, OH, H or CH_3), Trp, Cys, or
 39
                B-Nal;
      A<sup>7</sup> =
 40
                1-methyl-His, 3-methyl-His, or His;
      provided that if A^0 is present, A^1 cannot be pGlu;
      and if A^0 or A^1 is present, A^2 cannot be pGlu; and
 42
      when {\bf A}^0 is deleted and {\bf A}^1 is pGlu, {\bf R}_1 must be H
 43
      and R2 must be the portion of Glu that forms the imine
      ring in pGlu; and further provided that W can be:
 45
46
               Z1
-NH-CH-R2-C-V
47
48
      wherein R_3 is CHR_{14}-(CH_2)<sub>n1</sub> (where R_{14} is
49
      either H or OH; and n1 may be either of 1 or 0), or is
50
      deleted, and \mathbf{Z}_1 is the identifying group of any one of
51
      the amino acids Gly, Ala, cyclohexyl-Ala, Val, Leu, Ile,
52
     Ser, Asp, Asn, Glu, Gln, p-X-Phe (where X = H, F, Cl,
53
     Br, NO_2, OH, or CH_3), Trp, Cys, Met, Pro, HyPro, or
54
     isopropyl, cyclohexylmethyl, ß-nal, ß-napthylmethyl, or
55
     phenylmethyl; and V is either
56
     OR_4, or
57
58
     where R_4 is any of C_{1-20} alkyl, C_{3-20} alkenyl,
59
60
     C_{3-20} alkinyl, phenyl, napthyl, or C_{7-10}
61
     phenylalkyl, and each R_5, and R_6, independently, is
62
     any of H, C_{1-12} alkyl, C_{7-10} phenylalkyl, lower
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63
  64
  65
       where R_{15} is any cf H, C_{1-12} alkyl, C_{7-10}
  66
       phenylalkyl, or lower acyl; provided that when one of
  68
       R_5 or R_6 is NHR_{15}, the other is H; and provided
       that any asymmetric carbon atom can be R, S or a racemic
  69
  70
       mixture; and further provided that each R_1 and R_2,
  71
       independently, is H, C_{1-12} alkyl, C_{7-10} phenylalkyl,
       COE_1 (where E_1 is C_{1-20} alkyl, C_{3-20} alkenyl,
       C_{3-20} alkinyl, phenyl, naphthyl, or C_{7-10}
 73
       phenylalkyl), or lower acyl, and \mathbf{R}_{1} and \mathbf{R}_{2} are
       bonded to the N-terminal amino acid of said peptide, and
 75
       further provided that when one of R_1 or R_2 is
 76
       \mathtt{COE}_1, the other must be H, or a pharmaceutically
 77
       acceptable salt thereof.
 78
 1
                 5. The therapeutic peptide of claim 4 wherein
       A^0 = Gly, D-Phe, or is deleted;
 2
      A^1 = p-Glu, D-Phe, D-Ala, D-G-Nal, D-Cpa, or D-Asn;
      A^2 = Gln, His, 1-methyl-His, or 3-methyl-His;
      A^4 = Ala;
5
      A^5 = Val;
      A^6 = Sar, Gly, D-Phe, or D-Ala;
8
      provided that where R_3 is CH_2-CH_2, Z_1 is the
9
      identifying group of Leu or Phe; or where R3 is CH2,
10
11
      \mathbf{Z}_1 is the identifying group of \mathbf{B}	ext{-Leu} or Leu; or where
      \mathbf{R_3} is CHOH-CH<sub>2</sub>, \mathbf{Z_1} is the identifying group of Leu
12
13
      or is isopropyl, cyclohexylmethyl, ß-naphthylmethyl, or
      phenylmethyl; provided that where V is
14
14
16
17
      each R_5 and R_6 is H.
```

_ 49_ _

1 The therapeutic peptide of claim 5 wherein 2 V is NHR_6 and R_6 is H.

1 The therapeutic peptide of claim 5 of the 2

formula:

3

1

2

3

6

7

8

9

3 pGlu-Gln-Trp-Ala-Val-Gly-His-statine-amide.

1 The therapeutic peptide of claim 5 of the 2 formula:

 ${\tt D-p-Cl-Phe-Gln-Trp-Ala-Val-Gly-His-G-Leu-NH}_2\;.$

A therapeutic peptide comprising between eight and ten amino acid residues, inclusive, said peptide being an analog of one of the following naturally occurring peptides terminating at the carboxy-terminus with a Met residue: (a) litorin; (b) neuromedin; (c) the ten amino acid carboxy-terminal region of mammalian gastrin releasing peptide; and (d) the ten amino acid carboxy-terminal region of amphibian bombesin, said therapeutic peptide being of the formula:

10
$$R_1$$
 $A^0-A^1-A^2-Trp-A^4-A^5-A^6-A^7-W$
 R_2

13 wherein

 $A^0 =$ 14 Gly, Nle, α -aminobutyric acid, or the D-isomer 15 of any of Ala, Val, Gln, Asn, Leu, Ile, Met, 16 p-X-Phe (where X = F, Cl, Br, NO_2 , OH, H or 17 CH3), Trp, Cys, or B-Nal, or is deleted;

_ 50 _

```
\lambda^1 =
                 the D- or L-isomer of any of pGlu, Nle, or
  19
                 \alpha-aminobutyric acid, or the D-isomer of any of
  20
                 Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe (where
  21
                 X = F, Cl. 3r, NO_2, OH, H or CH_3), Trp, Cys.
  22
                 or G-Nal, or is deleted;
  23
                 pGlu, Gly, Ala, Val, Gln, Asn, Leu, Ile, Met,
                 p-X-Phe (where X = F, Cl, Br, NO<sub>2</sub>, OH, H or )
  25
                 CH3), Trp, Cys, B-Nal, His, 1-methyl-His, or
 26
                 3-methyl-His;
 27
                 Ala, Val, Gln, Asn, Gly, Leu, Ile, Nle,
 28
                 \alpha-aminobutyric acid, Met, p-X-Phe (where X = F,
 29
                 Cl, Br, NO<sub>2</sub>, OH, H or CH<sub>3</sub>), Trp, Cys, or
 30
                B-Nal;
      \lambda^5 =
 31
                Gln, Asn. Gly, Ala, Leu, Ile, Nle,
 32
                \alpha-aminobutyric acid, Met, Val, p-X-Phe (where X
 33
                = F, Cl, Br, OH, H or CH<sub>3</sub>), Trp, Thr, or B-Nal;
34
                Sar, Gly, or the D-isomer of any of Ala, Val,
35
                Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F,
36
                C1, Br, NO_2, OH, H or CH_3), Trp, Cys, or
37
                B-Nal;
38
                1-methyl-His, 3-methyl-His, or His;
      provided that if A^0 is present, A^1 cannot be pGlu; and
39
      if A^0 or A^1 is present, A^2 cannot be pGlu; and when
      {\tt A}^0 is deleted and {\tt A}^1 is pGlu, {\tt R}_1 must be H and {\tt R}_2
41
     must be the portion of Glu that forms the imine ring in
43
     pGlu; and further provided that W can be:
```

wherein R₄ is CH₂-NH, CH₂-S, CO-CH₂, or CH₂-CH₂ and each as T

 47 CH₂-CH₂, and each of 2 1 and 2 2, independently, is

the identifying group of any one of the amino acids Gly,
Ala. Val. Lev. Tip. Gr.

9 Ala, Val, Leu, Ile, Ser, Asp, Asn, Glu, Gln, 3-Nai,

81

_ 51 _

```
50 p-X-Phe (where X = H, F, Cl, Br, NO_2, OH or CH_3), Trp,
        Cys, Met, Pro, HyPro, cyclohexyl-Ala, or cyclohexylmethyl;
   52
        provided that where \mathbf{R}_4 is \mathbf{CH}_2\mathbf{-NH} and \mathbf{Z}_2 is the
   53
        identifying group of any one of the amino acids Gly, Ala,
   54
        Val, Leu,
   55
        Ile, Ser, Asp, Asn, Glu, Gln, p-X-Phe (where X = H, F,
  56
       C1, Br, NO_2, OH or CH_3), Trp, Cys, Met, Pro, HyPro,
  57
       or cyclohexylmethyl, Z_1 can only be the identifying
  58
       group of any one of the amino acids Ser, Asp, Glu, Cys,
  59
       Pro, HyPro, or cylcohexylmethyl; and provided that where
  60
       R_4 is CH_2-NH and Z_1 is the identifying group of
       any one of the amino acids Gly, Ala, Val, Leu, Ile, Ser,
  61
  62
       Asp, Asn, Glu, Gln, p-X-Phe (where X = H, F, Cl, Br,
  63
       \mathrm{NO}_2, OH or \mathrm{CH}_3), Trp, Cys, Met, Pro, or HyPro, \mathrm{Z}_2
 64
       can only be the identifying group of any one of the
 65
       amino acids Ser, Asp, Glu, Cys, Pro, HyPro, or
 66
       cylcohexylmethyl; and V is either OR_5 or
 67
 68
 69
 70
      where each R_8, R_5, R_6, and R_7, independently, is
 71
      H, lower alkyl, lower phenylalkyl, or lower
72
      naphthylalkyl; and provided that any asymmetric carbon
73
      atom can be R, S or a racemic mixture; and further
74
      provided that each R_1 and R_2, independently, is H,
75
      ^{\rm C}_{\rm 1-12} alkyl, ^{\rm C}_{\rm 7-10} phenylalkyl, ^{\rm COE}_{\rm 1} (where ^{\rm E}_{\rm 1}
76
      is C_{1-20} alkyl, C_{3-20} alkenyl, C_{3-20} alkinyl,
77
      phenyl, naphthyl, or C<sub>7-10</sub> phenylalkyl), or lower
78
      acyl, and \mathbf{R}_1 and \mathbf{R}_2 are bonded to the N-terminal
79
      amino acid of said peptide, and further provided that
80
      when one of R_1 or R_2 is COE_1, the other must be H,
      or a pharmaceutically acceptable salt thereof.
```

- 52 **-**

```
1 10. The therapeutic peptide of claim 9 wherein A^0 = Gly, D-Phe, or is deleted;
```

- 3 $A^1 = p-Glu$, D-Phe, D-Ala, D- β -Nal, D-Cpa, or D-Asn;
- 4 $A^2 = Gln$, His, 1-methyl-His, or 3-methyl-His;
- 5 $A^4 = Ala;$
- 6 $A^5 = Val;$
- 7 $A^6 = Sar, Gly, D-Phe, or D-Ala;$
- 8 $A^7 = His;$
- where R_4 is CH_2 -NH, each Z_1 is cyclohexylmethyl or
- is the identifying group of Leu or Phe; or Z_2 is the
- 11 identifying group of Met, Leu or Phe.
- 1 11. The therapeutic peptide of claim 9 wherein
 - A^{1} is D-B-Nal, each of Z_{1} and Z_{2} , independently,
 - 3 is Leu or Phe.
 - 1 12. The therapeutic peptide of claim 11 of the
 - 2 formula:
 - 3 $D-B-Nal-Gln-Trp-Ala-Val-Gly-His-Leu\psi[CH_2NH]Leu-NH_2$.
 - 1 13. The therapeutic peptide of claim 11 of the
 - 2 formula:
 - 3 D-β-Nal-Gln-Trp-Ala-Val-Gly-His-Leuψ[CH₂NH]Phe-NH₂.
 - 1 14. The therapeutic peptide of claim 9 wherein
 - 2 R_4 is CH_2 -NH, and said carbon atom bonded to Z_2 is
 - of said R configuration.
 - 1 15. The therapeutic peptide of claim 14 of the
 - 2 formula
 - D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu ψ [CH₂NH]-D-Phe-NH₂.
 - 1 16. The therapeutic peptide of claim 9 where
 - 2 A^0 or A^1 is a D amino acid and V is OR_4 .

23

24

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27

```
1
                17. The therapeutic peptide of claim 16 of the
  2
      formula:
  3
             D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu-Met-methylester.
 1
                     A therapeutic peptide comprising between
 2
      seven and nine amino acid residues, inclusive, said
 3
      peptide being an analog of one of the following
 4
      naturally occurring peptides terminating at the
      carboxy-terminus with a Met residue: (a) litorin; (b)
 5
      neuromedin; (c) the ten amino acid carboxy-terminal
 6
      region of mammalian gastrin releasing peptide; and (d)
 8
      the ten amino acid carboxy-terminal region of amphibian
     bombesin, said therapeutic peptide being of the formula:
 10
             A^0 - A^1 - A^2 - Trp - A^4 - A^5 - A^6 - A^7 - W
11
12
13
     wherein
     A^0 =
14
               Gly, Nle, \alpha-aminobutyric acid, or the
15
               D-isomer of any of Ala, Val, Gln, Asn, Leu,
16
               Ile, Met, p-X-Phe (where X = F, Cl, Br, NO_2,
17
               OH, H or CH_3), Trp, Cys, or B-Nal, or is
18
               deleted;
     A^1 =
19
               the D- or L-isomer of any of pGlu, Nle, or
20
              \alpha-aminobutyric acid, or the D-isomer of any
21
               of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe
22
               (where X = F, C1, Br, NO<sub>2</sub>, OH, H or CH<sub>3</sub>),
              Trp, Cys, or S-Nal, or is deleted;
     <sup>2</sup> =
              pGlu, Gly, Ala, Val, Gln, Asn, Leu, Ile, Met,
              p-X-Phe (where X = F, C1, Br, NO_2, OH, H or
```

CH₃), Trp, Cys, ß-Nal, His, l-methyl-His, or

3-methyl-His;

_ 54. _

Ala, Val. Glr. Asn. Gly. Leu. Ile. Nle. 29 α -aminobutyric acid, Met, p-X-Phe (where X = 30 F, C1, Br, NO_2 , OH, H or CH_3), Trp, Cys, or 31 B-Nal; 32 Gln, Asn, Gly, Ala, Leu, Ile, Nle, 33 α -aminobutyric acid, Met, Val, p-X-Phe (where 34 X = F, C1, Br, OH, H or CH_3), Trp, Thr, or 35 B-Nal; 36 Sar, Gly, or the D-isomer of any of Ala, Val, 37 Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F, 38 C1, Br, NO_2 , OH, H or CH_3), Trp, Cys, or-39 B-Nal: 40 1-methyl-His, 3-methyl-His, or His; provided that if A is present, A cannot be pGlu; 41 and if A^0 or A^1 is present, A^2 cannot be pGlu; and when ${\bf A}^0$ is deleted and ${\bf A}^1$ is pGlu, ${\bf R}_1$ must be H and R_2 must be the portion of Glu that forms the imine 44 45 ring in pGlu; and further provided that W can be:

49 wherein Z_1 is the identifying group of any one of the 50 amino acids Gly, Ala, Val, Leu, Ile, Ser, Asp, Asn, Glu, 51 Gln, p-X-Phe (where X = H, F, Cl, Br, NO_2 , OH or 52 CH3), Trp, Cys, Met, Pro, or HyPro; and each R9, 53 R_{10} , and R_{11} , independently, is H, lower alkyl, 54 lower phenylalkyl, or lower naphthylalkyl; and provided that any asymmetric carbon atom can be R, S or a racemic 56 mixture; and further provided that each R_1 and R_2 , 57 independently, is H, C_{1-12} alkyl, C_{7-10} phenylalkyl, 58 COE_1 (where E_1 is C_{1-20} alkyl, C_{3-20} alkenyl, 59 C_{3-20} alkinyl, phenyl, naphthyl, or C_{7-10}

WO 90/03980

1

2

3

1

2

3 -

- 55 -

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60
      phenylalkyl), cr lower acyl, and R_1 and R_2 are
 61
      bonded to the N-terminal amino acid of said peptide, and
      further provided that when one of R_1 or R_2 is
 63
      \mathtt{COE}_1, the other must be H, or a pharmaceutically
 64
      acceptable salt thereof.
 1
                     The therapeutic peptide of claim 18 wherein
      \lambda^0 = Gly, D-Phe, or is deleted;
 2
     A^{1} = p-Glu, D-Phe, D-Ala, D-B-Nal, D-Cpa, or D-Asn;
 3
     \lambda^2 = Gln, His, 1-methyl-His, or 3-methyl-His;
     A^4 = Ala;
     \lambda^5 = Val;
6
     A^6 = Sar, Gly, D-Phe, or D-Ala;
     A^7 = His;
     provided that \mathbf{Z}_1 is the identifying group of any one
10
     of the amino acids Leu or D or L p-X-Phe (where X = H,
11
     F, Cl. Br, NO_2, OH or CH_3); and each R_9, R_{10} and
12
     R<sub>11</sub>, independently, is H, lower alkyl, lower
13
     phenylalkyl, or lower naphthylalkyl.
1
                    The therapeutic peptide of claim 19
2
     wherein Z_1 is Leu, R_9 is H, and each R_{10} and R_{11}
     is lower alkyl.
                    The therapeutic peptide of claim 20 of the
     formula:
              D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu-ethylamide.
                   The therapeutic peptide of claim 20 of the
              22.
    formula:
              {\tt D-Phe-Gln-Trp-Ala-Val-Gly-His-Leu-NH}_2\;.
```

- 56 **-**

```
1
              23. A therapeutic peptide comprising between
2
     six and eight amino acid residues, inclusive, said
3
     peptide being an analog of one of the following
     naturally occurring peptides terminating at the
5
     carboxy-terminus with a Met residue: (a) litorin; (b)
6
     neuromedin; (c) the ten amino acid carboxy-terminal
7
     region of mammalian gastrin releasing peptide; and (d)
8
     the ten amino acid carboxy-terminal region of amphibian
9
     bombesin, said therapeutic peptide being of the formula:
```

13 wherein 14 Gly, Nle, α -aminobutyric acid, or the 15 D-isomer of any of Ala, Val, Gln, Asn, Leu, -16 Ile, Met, p-X-Phe (where X = F, Cl, Br, NO₂, 17 OH, H or CH_3), Trp, Cys, or β -Nal, or is 18 deleted: 19 the D- or L-isomer of any of pGlu, Nle, or 20 α -aminobutyric acid, or the D-isomer of any 21 of Ala, Val, Gln, Asn, Leu, Ile, Met, p-X-Phe 22 (where X = F, C1, Br, NO₂, OH, H or CH₃), 23 Trp, Cys, or B-Nal, or is deleted; 24 pGlu, Gly, Ala, Val, Gln, Asn, Leu, Ile, Met, 25 p-X-Phe (where X = F, Cl, Br, NO_2 , OH, H or 26 CH3), Trp, Cys, ß-Nal, His, 1-methyl-His, or 27 3-methyl-His; 28 Ala, Val, Gln, Asn, Gly, Leu, Ile, Nle, 29 α -aminobutyric acid, Met, p-X-Phe (where X = 30 F, Cl, Br, NO_2 , OH, H or CH_3), Trp, Cys, or 31 B-Nal;

```
- 57 -
                Gln. Asn. Gly. Ala. Leu. Ile. Nle.
33
                \alpha\text{--aminobutyric} acid, Met, Val, p-X-Phe (where
34
                X = F, C1, Br, OH, H or CH_3), Trp, Thr, or
35
                B-Nal;
36
                Sar, Gly, or the D-isomer of any of Ala, Val,
37
                Gln, Asn, Leu, Ile, Met, p-X-Phe (where X = F,
38
                Cl, Br, NO<sub>2</sub>, OH, H or CH<sub>3</sub>), Trp, Cys, or
39
                B-Nal;
     A^7 =
40
                l-methyl-His, 3-methyl-His, or His;
     provided that if A^0 is present, A^1 cannot be pGlu;
     and if A<sup>0</sup> or A<sup>1</sup> is present, A<sup>2</sup> cannot be pGlu; and
     when A^0 is deleted and A^1 is pGlu, R_1 must be H
     and R2 must be the portion of Glu that forms the imine
44
     ring in pGlu; and further provided that W can be:
45
```

49 wherein each R_{12} and R_{13} , independently, is H, lower alkyl, lower phenylalkyl, or lower naphthylalkyl; 50 provided that any asymmetric carbon atom can be R, S or 51 a racemic mixture; and further provided that each R_1 52 and R_2 , independently, is H, C_{1-12} alkyl, C_{7-10} 53 phenylalkyl, COE_1 (where E_1 is C_{1-20} alkyl, 54 55 C_{3-20} alkenyl, C_{3-20} alkinyl, phenyl, naphthyl, or 56 C_{7-10} phenylalkyl), or lower acyl, and R_1 and R_2 57 are bonded to the N-terminal amino acid of said peptide, and further provided that when one of R_1 or R_2 is 58 59 \mathtt{COE}_1 , the other must be H, or a pharmaceutically 60 acceptable salt thereof.

- 58 -

```
24. The therapeutic peptide of claim 23 wherein
      A^0 = Gly, D-Phe, or is deleted;
 2
      A^1 = p-Glu, D-Phe, D-Ala, D-S-Nal, D-Cpa, or D-Asn;
      A^2 = Gln, His, 1-methyl-His, or 3-methyl-His;
      A^4 = Ala;
 5
      A^5 = Val;
      A<sup>6</sup> = Sar, Gly, D-Phe, or D-Ala;
      A^7 = His;
      where each R_{12} and R_{13}, is H; and each R_1 and
      R<sub>2</sub>, independently, is H, lower alkyl, or lower acyl.
 Ŀ
                25. The therapeutic peptide of claim 24
     wherein either of N _{12} or N _{13} is other than H, \rm A^7 must be His, \rm A^6 must be Gly, \rm A^5 must be Val, \rm A^4
 2
 3
     must be Ala, and A<sup>2</sup> must be His.
4
1
                26. The therapeutic peptide of claim 24
     wherein either of R_1 or R_2 is other than H, A^1
3
     must not be deleted.
1
                27. The therapeutic peptide of claim 4, 9, 18,
     or 23 wherein said analog is at least 25% homologous
2
     with said naturally occurring peptide.
1
                     The therapeutic peptide of claim 27
     wherein said analog is at least 50% homologous with said
     naturally occurring peptide.
1
               29. A bombesin therapeutic peptide of the
2
     formula:
```

pGlu-Gln-Arg-Leu-Gly-Asn-Gln-Trp-Ala-Val-Gly-His-

3

Statine.

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- 59 -
```

```
1
                 30. An effective bombesin antagonistic peptide
 2
      containing the amino acid formula:
             A<sup>1</sup>-A<sup>2</sup>-A<sup>3</sup>-A<sup>4</sup>-A<sup>5</sup>-A<sup>6</sup>-A<sup>7</sup>-A<sup>8</sup>-A<sup>9</sup>-A<sup>10</sup>-
 3
 5
              A<sup>11</sup>-A<sup>12</sup>-A<sup>13</sup>-A<sup>14</sup>-N

-
 6
 7
8
9
      wherein
10
                 pGlu, D or L, or is deleted;
                 Gln, Asn, Gly, Ala, Leu, Ile, Nle,
12
                 \alpha-aminobutyric acid, Met, Val, Phe, p-X-Phe
13
                 (X = F, Cl, Br, OH or CH<sub>3</sub>), Trp,
14
                 B-naphthylalanine or is deleted;
15
                Arg, D-Arg, Lys, D-Lys or is deleted;
                Gln, Asn, Gly, Ala, Leu, Ile, Nle,
17
                \alpha-aminobutyric acid, Met, Val, Phe, p-X-Phe
18
                 (X = F, Cl, Br, OH or CH<sub>3</sub>), Trp,
19
                \beta-naphthylalanine or is deleted ;
20
                Gln, Asn, Gly, Ala, Leu, Ile, Nle,
21
                \alpha-aminobutyric acid, Met, Val, Phe, D-Phe,
22
                p-X-Phe (X = F, Cl, Br, OH or CH<sub>3</sub>), Trp,
23
                B-naphthylalanine, D-Ala or is deleted;
24
                Gln, Asn, Gly, Ala, D-Ala, N-Ac-D-Ala, Leu,
25
                Ile, Nle, \alpha-aminobutyric acid, Met, Val, Phe,
26
                p-X-Phe (X = F, Cl, Br, OH or CH<sub>3</sub>), Trp,
27
                p-Glu, ß-naphthylalanine or is deleted;
28
                Gln, Asn, Gly, Ala, Leu, Ile, Nle,
29
                \alpha-aminobutyric acid, Met, Val, Phe, D-Phe,
30
                p-X-Phe(X = F, Cl, Br, OH or CH_3), Trp, Lys,
31
                His, or \(\beta\)-naphthylalanine;
32
                Trp or Met;
```

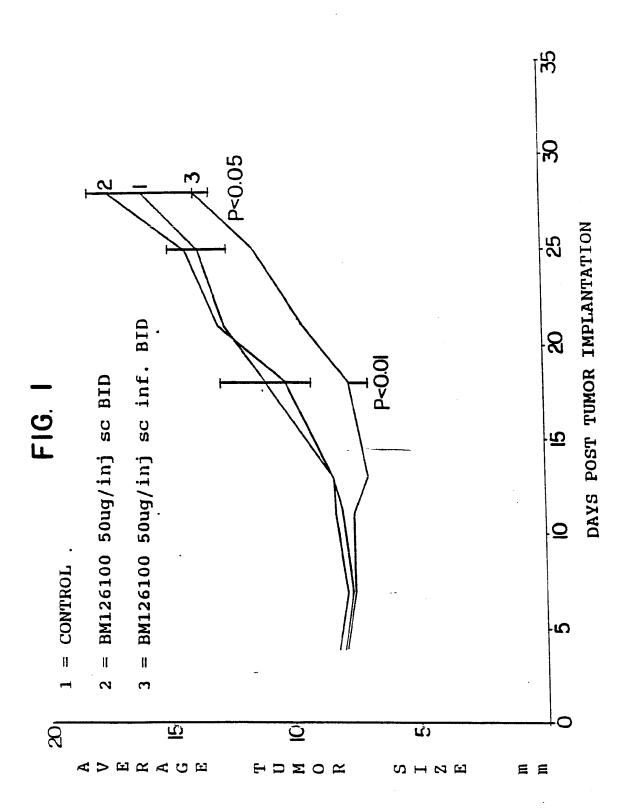
- 60 -

```
33
                 Gln, Asn, Gly, Ala, Leu, Ile, Nle,
  34
                 \alpha-aminobutyric acid, Met, Val, Phe, p-X-Phe
 35
                 (X = F, C1, Br, OH or CH<sub>3</sub>), Trp, or
 36
                 \beta-naphthylalanine, D or L;
       A<sup>10</sup>
 37
                 Gln, Asn, Gly, Ala, Leu, Ile, Nle,
 38
                 lpha-aminobutyric acid, Met, Val, Phe, p-X-Phe
 39
                 (X = F, C1, Br, OH or CH<sub>3</sub>), Trp, Thr, or
 40
                 ß-naphthylalanine;
      A^{11} =
 41
                 Gly, Phe, D or L;
      A^{12} =
 42
                 His, Phe, or p-X-Phe (X = F, Cl, Br, OH,
 43
                 CH<sub>3</sub>), D or L;
 44
                 Gln, Asn, Gly, Ala, Leu, Ile, Nle,
 45
                 \alpha-aminobutyric acid, Met, Val, Phe, p-X-Phe
 46
                 (X = F, C1, Br, OH or CH<sub>3</sub>), Trp, or
 47
                ß-naphthylalanine;
 48
                Gln, Asn, Gly, Ala, Leu, Ile, Nle,
49
                \alpha-aminobutyric acid, Met, Val, Phe, p-X-Phe
50
                (X = F, Cl, Br, OH or CH<sub>3</sub>), Trp, or
. 51
                ß-naphthylalanine;
52
      provided that
53
                each R_1, R_2, R_3, and R_4, independently,
54
      is H, C_{1-12} alkyl, C_{7-10} phenylalkyl, COE_1 (where
55
      E_1 is C_{1-20} alkyl, C_{3-20} alkenyl, C_{3-20} alkinyl,
      phenyl, naphthyl, or C_{7-10} phenylalkyl), or COOE_2
56
• 57
      (where E_2 is C_{1-10} alkyl or C_{7-10} phenylalkyl),
58
      and R_1 and R_2 are bonded to the N-terminal amino
59
      acid of said peptide, which can be A<sup>1</sup>, A<sup>2</sup>, A<sup>3</sup>,
     A^4, A^5, A^6, or A^7, and further provided that
60
61
     when one of R_1 or R_2 is COE_1 or COOE_2, the other
     must be H, and when one of R_3 or R_4 is COE_1 or
62
     COOE2, the other must be H, and further provided that
63
     when A^1 = pGlu, R_1 must be H and R_2 must be the
64
     portion of Glu that forms the imine ring in pGlu; and
     for each of the residues A^7, A^8, A^9, A^{11}, A^{12},
66
```

WO 90/03980

- 61 -

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and {\mbox{A}}^{13}, independently, the carbon atom participating
 67
 68
       in the amide bond between that residue and the nitrogen
 69
       atom of the alpha amino group of the adjacent amino acid
70
      residue may be a carbonyl carbon or may be reduced to a
71
      methylene carbon, provided that at least one such carbon
72
      atom must be reduced to a methylene carbon,
73
                said peptide further comprising
                                                        ()
      A^5 = cys;
74
      A^6 = Cys or a D-isomer of any of said amino acids;
75
      A^7 = pGlu, Cys, 1-methyl-His, or 3-methyl-His;
76
      A^9 = Cys;
77
      A^{11} = Sar, or the D-isomer of any of Ala, Val, Gln,
78
79
      Asn, Leu, Ile, Met, p-X-Phe (where X = F, Cl, Br, NO<sub>2</sub>,
80
      OH, or CH3), Trp, Cys, or B-Nal;
      A^{12} = 1-methyl-His, or 3-methyl-His;
81
      and where A<sup>14</sup> may be deleted.
82
83
84
                     The therapeutic peptide of claim 30 of the
85
      formula:
86
      D-p-Cl-Phe-Gln-Trp-Ala-Val-Gly-His-Leuw[CH2NH]Phe-NH2
```



SUBSTITUTE SHEET

Litorin

A1 A2 A3 A4 A5 A6 A7 A8 A9 pGlu-Gln-Trp-Ala-Val-Gly-His-Phe-Met

Neuromedin C

A0 A1 A2 A3 A4 A5 A6 A7 A8 A9 Gly-Ser-His-Trp-Ala-Val-Gly-His-Leu-Met

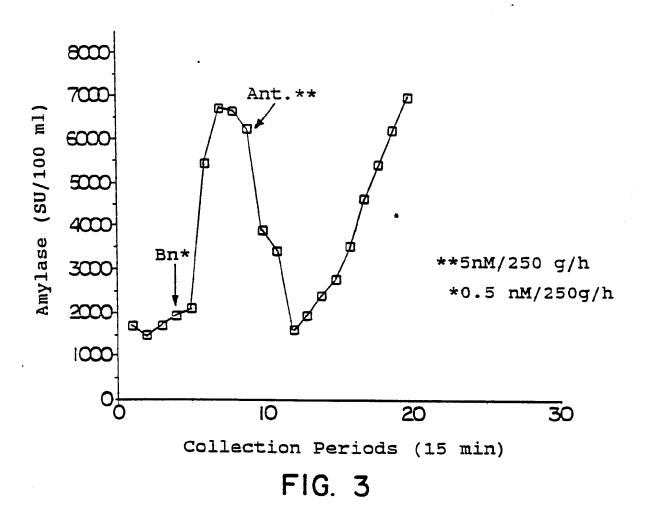
Bombesin (last 10 amino acids)

A0 A1 A2 A3 A4 A5 A6 A7 A8 A9 Gly-Asn-Gln-Trp-Ala-Val-Gly-His-Leu-Met

human GRP (last 10 amino acids)

A0 A1 A2 A3 A4 A5 A6 A7 A8 A9 Gly-Asn-His-Trp-Ala-Val-Gly-His-Leu-Met

FIG. 2



INTERNATIONAL SEARCH REPORT

International Application No. PCT/US89/04616

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TDC (A)	io internat	ional Patent Classification (IPC) or to both Natio X 7/02,7/06,7/08,7/10,7/30	nai Classification and IPC	
			227 220 220	
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" FIELD:	3 SEARCE	IED		
		Minimum Document	ation Searched 7	
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		to the strain distribution additional to	ire includes in the Fields Searched	
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III. BOCL	MENTS C	ONSIDERED TO BE RELEVANT		
Category *	Citat	ion of Document, 11 with indication, where appro	opnate, of the relevant passages 12	Relevant to Claim No. 13
				Herevalle to Claim Rd. 13
x	US. Z	4,207,311 (Brown et	al.),	9
	10	Tune 1980. See column	2. line 29 and	
		l in particular.		
	Clair	i i ili parcicatar.		
x	ric 7	A, 4,331,661 (Marki et	al.).	9
Λ.	03,	May 1982. See the abs	tract in par-	-
			crace in par-	
	ticu:	Lar.		
				9
X	US,	A, 4,613,586 (Barchas	et al.),	ا ع
	23,	September 1986. See t	able 1, lines	
	8-10	in particular.		
P, A	us,	A, 4,803,261 (Coy et a	11.),	1-31
- 1		February 1989.		
X	Δm .T	. of Physiol, (Marylar	nd, USA), issued	1-3
**	1005	, (Heinz-Erian et al.)	. "[D-phe ¹²] bom-	
	1900	n analogues: a new cl	ass of hombesin	
	pesi	n anatogues: a new ca	25 C 439-G442	i .
	rece	ptor antagonists", pag	Jes G 439-G442.	
	See	the abstract in partic	cutat.	
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* 5pecis	M categories	of cited documents: 19	"T" later document mubished after or prigrity date and not in conf	the international filing date
con	sument deri	ning the general state of the art which is not be of particular relevance	cited to understand the princip	
"E" carl	ier docume	nt but published on or after the international	invention "X" document of particular releval	and the demand a contra
	ig date	the manufacture of the second	cannot be considered novel o	r cannot be considered to
whi	CR IS CITED	th may throw doubts on priority claim(s) or to establish the publication date of another	"Y" document of particular relevan	حديده عمستمام مراه بمعد
		rring to an oral disclosure, use, exhibition or	cannot be considered to involve	BU IUAGUIAS ELSO WJOG JE
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			"4" document member of the same	patent family
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FURTHE	R INFORMATION CONTINUED FROM THE SECOND SHEET	1/ 0309/ 04016
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х	Proc. Natl. Acad. Sci. USA (Washington, D.C., USA) volume 82, issued November, 1985. (Zachary et al.), "High-affinity receptors for peptides of the bombesin family in Swiss 3T3 cells", pages 7616-7620. See Table 2, Neuromedin C and B in particular.	9
	•	
V. 🗌 089	SERVATIONS WHERE CERTAIN CLAIMS WERE FOUND UNSEARCHABLE!	···
This intern	ational search report has not been established in respect of certain claims under Article 17(2) (a) for	the following reasons:
1. Clain	numbers because they relate to subject matter 13 not required to be searched by this Auth	ority, namely:
	•	
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2. Claim ments	numbers because they relate to parts of the international application that do not comply wit s to such an extent that no meaningful international search can be carried out 13, specifically:	h the prescribed require-
	numbers because they are dependent claims not drafted in accordance with the second and fule 6.4(a).	third sentences of
VI. 🗌 088	ervations where unity of invention is lacking:	
This Interna	stional Searching Authority found multiple inventions in this international application as follows:	
1. As all	required additional search fees were timely paid by the applicant, this international search report coverintened applicants.	ee all engagements are
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thase	ly some of the required additional search fees were timely paid by the applicant, this international se claims of the international application for which fees were paid, specifically claims:	erch report covers anly
3. No req	wired additional search fees were timely paid by the applicant. Consequently, this international searce ention first mentioned in the claims; it is covered by claim numbers:	h report is restricted to
_	searchable claims could be searched without effort justifying an additional fee, the International Searchant of any additional fee.	Ching Author ty 2 2 2*
Remark on F		
	ditional search fees were accompanied by applicant's protest. test accompanied the payment of additional search fees.	
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PCT/US89/04616

Category •	MENTS CONSIDERED TO BE RELEVANT (CONTINUED FROM THE SECOND SHEE	T)
Category	Citation of Document, with indication, where appropriate, of the relevant passages	Relevant to Claim !
X Y	Cancer Surveys (Oxford, England) volume 4, no. 4, issued 1985 (Cuttitta et al.), "Autocrine growth factors in human small cell lung cancer", pages 707-727. See page 718 and Table 2 in particular.	1-3,9, U,3U-31
Y	J. Med. Chem. (Washington, D.C., USA) volume 30 issued 1987 (Sasaki et al.) "Solid-phase synthesis and biological Properties of \(\mathbb{C}(CH2NH) \) Pseudopeptide analogues of a highl Potent somatostatin octapeptide", pages 1162-1166. See pages 1162, 1164, 1166 in particular.	10,30-31 Y
X	Chemical Abstract, (Columbus, Ohio, USA) volume 109, issued 1988, (Coy et al.) "Probing peptide backbone function in bombesin. A reduced peptide bond analog with potent and specific receptor antagonist activity", the abstract No. 32216K, J. Biol. Chem. 1988, 263 (11), 5056-60 (Eng).	1-3,30
	Chemical Abstract, (Columbus, Ohio, USA) volume 109, issued 1988, (WQ11 et al.), "[Leu ¹³ + (CH ₂ NH) Leu ¹⁴] bombesin is a special bombesin receptor antagonist in Swiss 3T3 cells", the abstract no 163928s, Biochem. Biophys. Res. Commun. 1988, 155(1) 359-65(Eng).	1-3,30
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